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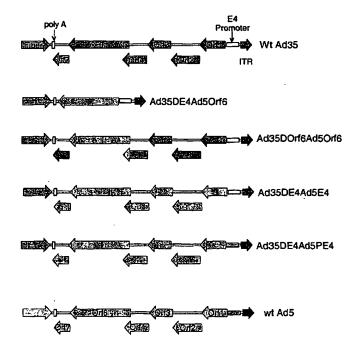
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[Continued on next page]

(54) Title: METHODS FOR PROPAGATING ADENOVIRUS AND VIRUS PRODUCED THEREBY



(57) Abstract: Various methods propagating and rescuing multiple serotypes of replication-defective adenovirus in a single adenoviral E1-complementing cell line are Typically, replication-defective disclosed. adenovirus vectors propagate only in cell lines which express E1 proteins of the same serotype or subgroup as the vector. The disclosed methods offer the ability to propagate vectors derived from multiple adenoviral serotypes in a single production cell line which expresses E1 proteins from a single serotype. Propagation in this manner is accomplished by providing all or a portion of an E4 region in cis within the genome of the replication-defective adenovirus. The added E4 region or portion thereof is cloned from a virus of the same or highly similar serotype as that of the E1 gene product(s) of the complementing cell line. Interaction between the expressed E1 of the cell line and the heterologous E4 of the replication-defective adenoviral vectors enables their propagation and rescue. The invention bypasses a need in the art to customize specific cell lines to the serotype or subgroup of the adenoviral vector being propagated and enables one to easily and rapidly develop alternative adenoviral serotypes as gene delivery vectors for use as vaccines or as a critical component in gene therapy.

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TITLE OF THE INVENTION

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METHODS FOR PROPAGATING ADENOVIRUS AND VIRUS PRODUCED THEREBY

CROSS-REFERENCE TO RELATED APPLICATIONS

The present application claims the benefit of application serial nos. 60/458,825, filed March 28, 2003; 60/455,312, filed March 17, 2003; 60/455,234, filed March 17, 2003; and 60/405,182, filed August 22, 2002.

FIELD OF THE INVENTION

The present invention concerns various methods to propagate and rescue multiple serotypes of replication-defective adenovirus in a single adenoviral E1-complementing cell line. Typically, replication-defective adenovirus vectors propagate only in cell lines which express E1 proteins of the same serotype or subgroup as the vector. The methods disclosed herein offer the ability to propagate vectors derived from multiple serotypes in a single cell line expressing E1 proteins from a single serotype. Such propagation of a wide range of vectors in one cell line is accomplished by providing all or a portion of an E4 region in cis within the genome of the replication-defective adenovirus. The added E4 region or portion thereof is cloned from a virus of the same or highly similar serotype as that of the E1 gene product(s) of the complementing cell line. Interaction between the E1 gene products of the cell line and the heterologous E4 gene products of the replication-defective adenoviral vector enables the propagation and rescue of the recombinant replication-defective adenovirus vectors. The invention, therefore, bypasses an existing need in the art to customize complementing cell lines to the specific serotype or subgroup of the adenoviral vector being propagated or, alternatively, to have to transfect a cell line with an E4 region and then regulate the expression in trans of the E4 region within the E1 complementing cell line.

BACKGROUND OF THE INVENTION

Beginning with the first human adenoviruses (Ads) isolated over four decades ago (Rowe et al., Proc. Soc. Exp. Biol. Med., 84:570-579, 1953), over 100 distinct serotypes of adenovirus have been isolated which infect various mammalian species, 51 of which are of human origin (Straus, Adenovirus infections in humans. In The Adenoviruses. 451-498, 1984; Hierholzer et al., J. Infect. Dis., 158: 804-813, 1988; Schnurr and Dondero, Intervirology., 36: 79-83, 1993; Jong et al., J Clin Microbiol., 37:3940-3945:1999). The human serotypes have been categorised into six subgenera (A-F) based on a number of biological, chemical, immunological and structural criteria; criteria which include hemagglutination properties of rat

and rhesus monkey erythrocytes, DNA homology, restriction enzyme cleavage patterns, percentage of G+C content and oncogenicity (Straus, Adenovirus infections in humans. In *The Adenoviruses*. 451-498, 1984; Horwitz, Adenoviridae and their replication, *In Virology*: 1679-172, 1990).

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Deletion of an essential E1 region common to the various adenovirus serotypes has enabled the use of adenovirus vectors as gene transfer vectors for vaccine and gene therapy purposes. Resultant replication-defective vectors are propagated in cell lines that provide the deleted E1 gene products in trans. Supplementation of the essential E1 gene products in trans in this manner works well when the E1 gene products are from the same or a highly similar serotype. As such, E1-deleted group C serotypes (Ad1, Ad2, Ad5 and Ad6) grow well in 293 or PER.C6 cells which contain and express the Ad5 E1 region. In contrast, E1-deleted serotypes other than group C, for example those from subgroups A, B, D, E, and F (e.g., Ad3, Ad4, and Ad7 to Ad51), do not replicate efficiently in 293 or PER.C6 cells. The Ad5 E1 sequences in 293 and PER.C6 cells do not fully complement the replication of these alternative serotypes. This presents a challenge due to the fact that the most characterized and studied complementing cell lines available for growth and propagation of adenovirus are based on E1 sequence from adenovirus serotype 5.

This inability to fully complement the replication of serotypes other than group C adenovirus in Ad5 E1 complementing cell lines has been attributed to the inability of Ad5 (group C) E1b 55K gene product to functionally interact with the E4 gene products of non-group C serotypes. While the interaction is conserved within members of the same subgroup, it is not well conserved between subgroups.

Hence, cell lines expressing both Ad5 E1 and ORF6 were generated and proved useful in complementing alternative adenovirus serotypes; see, e.g., Abrahamsen et al., 1997 J. Virol. 8946-8951. Such incorporation of E4 (or ORF6) into Ad 5 complementing cell lines as was done in Abrahamsen et al., supra, is known.

U.S. Patent No. 5,849,561 discloses complementation of an E1-deleted non-group C adenovirus vector in an Ad5-E1 complementing cell line which also expresses portions of the Ad5-E4 gene.

U.S. Patent No. 6,127,175, issued to Vigne, et al., discloses a stably transfected mammalian cell line which expresses a portion of the E4 region of adenovirus, preferably ORF6 or ORF6/7. Such a cell line is useful for complementation of recombinant Ad genomes deficient in the E4 region.

European Application EP 1 054 064 A1 discloses recombinant, replication deficient adenovirus 35 (Ad35) vectors and cell lines which complement in trans the growth of

these vectors. A cell line which expresses Ad5E1A and E2A genes (PER.C6) was shown to complement an Ad35-E1 deleted vector upon co-expression of Ad35-E1B proteins.

U.S. Patent No. 6,270,996, issued to Wilson, et al., discloses E1/E4 deleted adenovirus vectors and E1/E4(ORF6) cell lines which complement in trans virus growth without resulting in cell toxicity.

U.S. Patent No. 6,202,060, issued to Mehtali, et al., discloses adenoviral vectors wherein portions of the early genes are under control of an inducible promoter. The '060 patent also discloses complementing cell lines which may be used in tandem with these Ad vectors.

The generation of serotype-specific cell lines providing a complementing serotype-specific E1 gene product(s) in trans is known as well.

Although Ad5-based vectors have been used extensively in a number of gene therapy trials, there may be limitations on the use of Ad5 and other group C adenoviral vectors due to preexisting immunity in the general population due to natural infection. Ad5 and other group C members tend to be among the most seroprevalent serotypes. Immunity to existing vectors may develop as a result of exposure to the vector during treatment. These types of preexisting or developed immunity to seroprevalent gene delivery vectors may limit the effectiveness of gene therapy or vaccination efforts. Alternative adenovirus serotypes, thus, constitute very important targets in the pursuit of gene delivery systems capable of evading the host immune response.

There remains both a practical and commercial need for an adenovirus-based vaccine and/or gene therapy delivery system which allows for the production of multiple serotype recombinant adenovirus vectors in a single source complementing mammalian cell line. The present invention addresses and overcomes this deficiency in the art by disclosing novel methods for propagating multiple serotype recombinant Ad vectors in a single complementing cell line where the required serotype-specific sequences are provided *in cis*.

SUMMARY OF THE INVENTION

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The present invention relates to an enhanced means for propagating replication-defective adenovirus in an E1-complementing cell line(s) where the E1 gene product(s) being expressed is not native to the adenovirus being propagated. The method is based on Applicants' finding that supply, in cis, of a nucleic acid sequence encoding all or a portion of a heterologous adenoviral E4 region which is native to a virus of the same or highly similar serotype as the E1 gene product(s) of the complementing cell line enables the growth of adenoviral vectors of varying serotype in any single complementing cell line, despite the fact the cell line is not customized for the particular serotype of vector being propagated. This is of particular

importance given that existing and settled adenoviral E1-complementing cell lines (such as PER.C6™ and 293) are based on one of the most prominent adenovirus serotypes (Ad5) and are not suited for the large-scale propagation and rescue of alternative serotypes.

The basic steps involved in the propagation of adenoviral vectors in accordance with the methods of the instant invention are as follows: First, all or a portion of a heterologous adenoviral E4 region comprising nucleic acid sequence encoding at least open reading frame 6 (ORF6) is inserted into a replication-defective adenoviral vector. By "heterologous", Applicants mean that the nucleic acid sequence is not native to the viral vector being propagated, i.e., not normally present within a virus of the same or highly similar serotype. As will be described, the adenoviral E4 region or portion thereof can be either a nucleic acid sequence encoding ORF 6 or any larger portion of the E4 region, and includes nucleic acid comprising the complete E4 region with E4 promoter. The region into which the nucleic acid is incorporated is not limited, i.e., the insertion can be made into the complete E4 region with E4 promoter or into a smaller portion narrowing into the ORF6 region. Alternatively, the heterologous E4 region or portion thereof can be inserted into different areas of the genome such as the E1 or E3 regions. Further, the native E4 region or portion thereof can be deleted and replaced, or left intact. This is not deemed a critical element of the instant invention. What is a critical element is that the heterologous E4 region or portion thereof being inserted is native to a virus of the same or highly similar serotype as the E1 gene product(s) expressed by the complementing cell line.

Following the modification of the adenoviral vector of interest, the recombinant adenovirus is then introduced into an adenoviral E1-complementing cell line and allowed to propagate. The adenovirus is subsequently harvested and rescued from the complementing cell

The resultant virus can be studied and used in various gene therapy and vaccine line. efforts. The virus, therefore, forms an important aspect of the instant invention.

BRIEF DESCRIPTION OF THE DRAWINGS

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FIGURE 1 illustrates a transcription map for adenovirus serotype 5. The linear genome is divided into 100 map units as well as into r- and l- strands which designate the direction of transcription. Early transcription units are designated with an E and are active prior to viral DNA replication. Late transcription units are designated with and L and are active primarily after DNA replication. Promoters are represented as brackets and polyadenylation sites as arrowheads. The tripartite leader is designated 1, 2, and 3.

FIGURES 2A-1 through 2A-10 illustrate the nucleic acid sequence for the wildtype adenovirus 35 (SEQ ID NO: 1) utilized in the Examples.

FIGURE 3 illustrates the homologous recombination scheme utilized to recover pAd35 Δ E1.

FIGURE 4 illustrates the various configurations of the E4 regions (or portions) within the alternative serotype recombinants.

FIGURE 5 illustrates the homologous recombination scheme utilized to recover pAd35 Δ E1 Δ E4Ad5Orf6.

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FIGURE 6 illustrates the nucleic acid sequence encoding the gag expression cassette (SEQ ID NO: 2). The various regions of the figure are as follows: (1) a first underlined segment of nucleic acid sequence encoding the immediate early gene promoter region from human cytomegalovirus; (2) a first segment of lowercase letters which is not underlined, which segment of DNA contains a convenient restriction enzyme site; (3) a region in caps which contains the coding sequence of HIV-1 gag; (4) a second segment of lowercase letters which is not underlined, which segment of DNA contains a convenient restriction enzyme site; and (5) a second underlined segment, this segment containing nucleic acid sequence encoding a bovine growth hormone polyadenylation signal sequence.

FIGURE 7 illustrates the nucleic acid sequence encoding the SEAP expression cassette (SEQ ID NO: 3). The various regions of the figure are as follows: (1) a first underlined segment of nucleic acid sequence encoding the immediate early gene promoter region from human cytomegalovirus; (2) a first segment of lowercase letters which is not underlined, which segment of DNA contains a convenient restriction enzyme site; (3) a region in caps which contains the coding sequence of the human placental SEAP gene; (4) a second segment of lowercase letters which is not underlined, which segment of DNA contains a convenient restriction enzyme site; and (5) a second underlined segment, this segment containing nucleic acid sequence encoding a bovine growth hormone polyadenylation signal sequence.

FIGURE 8 illustrates *in vivo* expression of SEAP in C3H/HeN mice using 10¹⁰ vp doses of Ad35 vectors. This experiment was designed to address any effects of E3 deletion. The vectors were injected intramuscularly and the levels of SEAP expression were determined from the serum samples. Shown are geometric means for each cohort of 5 mice.

FIGURE 9 illustrates in vivo expression of SEAP in C3H/HeN mice using 10^10 vp doses of Ad35 vectors. This experiment was designed to address any effects of Ad5 sequence insertion into the Ad35 genome. The vectors were injected intramuscularly and the levels of SEAP expression were determined from the serum samples. Two extra cohorts received 10^10 vp and 10^9 vp of Ad5 vector. Shown are geometric means for each cohort of 5 mice.

FIGURES 10A-B illustrate *in vivo* SEAP expression using MRKAd5-based (A) and Ad35ΔE1ΔE4Ad5Orf6-based (B) vector in rhesus macaques. Shown are the serum antigen

levels for individual monkeys following a single intramuscular (i.m.) injection of 10¹¹ vp MRKAd5SEAP (filled circles), 10⁹ vp MRKAd5SEAP (open boxes) or 10¹¹ vp Ad35ΔE1SEAPΔE4Ad5Orf6.

FIGURE 11 illustrates *in vivo* SEAP expression in African green monkeys using

Ad5- and Ad35-based vectors. Shown are the antigen levels for each animal in serum samples collected two days after the treatment.

FIGURE 12 illustrates the homologous recombination scheme utilized to recover pAd24ΔΕ1.

FIGURE 13 illustrates the homologous recombination scheme utilized to recover NE1Ad5Orf6.

pAd24ΔE1Ad5Orf6.

FIGURE 14 illustrates the configuration of E4 regions in the Ad24 recombinants generated.

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FIGURE 15 illustrates the growth kinetics of the Ad24-based vectors in PER.C6 s.

rigures 16A-1 through 16A-10 illustrate the nucleic acid sequence for wild-type adenovirus serotype 24 (SEQ ID NO: 5). The ATCC product number for Ad24 is VR-259.

FIGURE 17 illustrates, in tabular format, gag-specific T cell responses in monkeys immunized with MRKAd5-HIVgag and Ad24 HIV vectors. Shown are the numbers of spot-forming cells per million PBMC following incubation in the absence (mock) or presence of Gag peptide pool. The pool consisted of 20-aa peptide overlapping by 10 aa and encompassing the entire gag sequence.

FIGURE 18 illustrates, in tabular format, the characterization of the gag-specific T cells in monkeys immunized with 10^11 vp of MRKAd5-HIV1gag and Ad24ΔE1gagΔOrf6Ad5Orf6. Shown are the percentages of CD3+ T cells that are either gag-specific CD4+ or gag-specific CD8+ cells. These values were corrected for mock values (<0.03%).

FIGURE 19 illustrates individual anti-p24 titers (in mMU/mL) in macaques immunized with gag-expressing adenovirus vectors.

FIGURE 20 illustrates *in vivo* expression of SEAP in C3H/HeN mice using 10¹⁰ vp doses of Ad24 vectors. The vectors were injected intramuscularly and the levels of SEAP expression were determined from the serum samples. Two extra cohorts received 10¹⁰ vp and 10⁹ vp of Ad5 vector. Shown are geometric means for each cohort of 5 mice.

FIGURE 21 illustrates in vivo SEAP expression using MRKAd5 and Ad24 vectors in rhesus macaques. Shown are the geometric means of the SEAP levels for cohorts of 3 monkeys. In bars are the standard errors of the geometric means.

FIGURE 22 illustrates a homologous recombination scheme to be utilized to recover pAd24 Δ E1 Δ E4Ad5Orf6.

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FIGURE 23 illustrates gag-specific T cell responses in rhesus macaques immunized following a heterologous Ad5/Ad6 prime-Ad24 boost regimen. a: Mock, no peptide: gag, 20-mer peptide pool encompassing entire gag sequence; b: Peak response after 2 or 3 doses of the priming vaccine; c: 3 wks prior to boost; d: 4 wks after boost; e: ND, not determined.

FIGURE 24 illustrates, in tabular format, the percentages of CD3⁺ T lymphocytes that are gag-specific CD8⁺ cells or gag-specific CD4⁺ cells determined after the Ad24 Boost Immunization (wk 60). Numbers reflect the percentages of circulating CD3+ lymphocytes that are either gag-specific CD4+ or gag-specific CD8+ cells. Mock values (equal to or less than 0.01%) have been subtracted.

FIGURE 25 illustrates gag-specific T cell responses in rhesus macaques immunized following a heterologous Ad 24 prime-Ad5 boost regimen. a: Mock, no peptide: gag, 20-mer peptide pool encompassing entire gag sequence; b: Peak response after 2 doses of the priming vaccine; c: Wk 24; d: 4 wks after boost; e: ND, not determined.

FIGURE 26 illustrates the homologous recombination scheme utilized to recover pAd34 Δ E1 Δ E4Ad5Orf6.

FIGURE 27 illustrates the homologous recombination scheme utilized to recover pMRKAd34 Δ E1 Δ E4Ad5Orf6.

FIGURES 28A-1 to 28A-9 illustrate a nucleic acid sequence for wild-type adenovirus serotype 34 (SEQ ID NO: 12). The ATCC product number for Ad34 is VR-716.

FIGURE 29 illustrates the time course of SEAP expression using MRKAd5 and Ad34 vectors in rhesus macaques. Data represent cohort geometric means.

FIGURE 30 illustrates, in tabular format, T cell responses induced using MRKAd5 and Ad34 vectors expressing HIV-1 gag. Data are expressed in numbers of spot-forming cells per million PBMC (SFC/10^6 PBMC). "a" refers to a 20-mer peptide pool with 10-aa overlap and encompassing the entire HIV-1 CAM1 gag.

FIGURE 31 illustrates, in tabular format, the levels of CD4+ and CD8+ Gag-specific T cells in Ad34-immunized macaques at week 12. "a" refers to a 20-mer peptide pool with 10-aa overlap and encompassing the entire HIV-1 CAM1 gag.

FIGURE 32 illustrates, in tabular format, T cell responses induced using a heterologous Ad34 prime/Ad35 boost regimen in macaques. "a" refers to a 20-mer peptide pool with 10-aa overlap and encompassing the entire HIV-1 CAM1 gag.

FIGURE 33 illustrates, in tabular format, the levels of CD4+ and CD8+ Gagspecific T cells in Ad34 primed/Ad35 boosted macaques at week 28. "a" refers to a 20-mer peptide pool with 10-aa overlap and encompassing the entire HIV-1 CAM1 gag.

DETAILED DESCRIPTION OF THE INVENTION

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The present invention details an efficient strategy for the propagation and rescue of alternative adenoviral serotypes utilizing available adenovirus production cell lines, nullifying the need to customize available cell lines for a specific serotype of interest. This is enabled by the incorporation of a critical E4 region into the adenovirus to be propagated.

The critical E4 region in the instant invention comprises, in the minimum, nucleic acid sequence encoding E4 ORF6 and can comprise the entire region of E4, inclusive of the promoter region. An important characteristic of the imported E4 region is that it is native to a virus of the same or highly similar serotype as the E1 gene product(s) (particularly E1B 55K) of the E1-complementing cell line, but heterologous to (i.e., non-native to a virus of the same serotype as) the adenoviral vector being propagated. As will be detailed below, the heterologous E4 region or portion thereof can be varied and can be inserted into the vector backbone at numerous locations.

The heterologous E4 region or portion thereof can, for instance, be a nucleic acid sequence encoding the entire open reading frame of the non-native E4. This segment of nucleic acid sequence can, in turn, be incorporated into the "native" entire E4 open reading frame of the recipient virus. In such an embodiment, the promoter native to the adenoviral vector would drive the expression of the non-native E4 region within the recombinant replication-defective adenoviral vector. Alternatively, the nucleic acid sequence encoding the entire open reading frame can be inserted into a different region of the adenoviral vector genome, such as for example the E1 or E3 regions. In this latter embodiment, the native E4 region or portion thereof can be deleted or left intact.

In another embodiment, the heterologous E4 region comprises a nucleic acid sequence encoding the entire open reading frame of E4 and includes a non-native E4 promoter. In this type of embodiment, the E4 region can be inserted into the location of the combined native E4 and E4 promoter region. The non-native E4 region in this embodiment would be driven by expression of the non-native E4 promoter. Alternatively, the nucleic acid sequence encoding the entire open reading frame and the non-native E4 promoter can be inserted into a different region of the adenoviral vector genome, such as for example the E1 or E3 regions. In this latter embodiment, the native E4 region or portion thereof can be deleted or left intact.

An alternative and further embodiment exists wherein the heterologous E4 region or portion thereof comprises nucleic acid sequence encoding a partial E4 region comprising ORF6 (one aspect of which is a region solely encoding ORF6). In this particular aspect of the invention, the heterologous non-native E4 protein can, in certain embodiments, replace the non-native ORF6 region or the entire E4-encoding region of the native virus. In the latter situation, the promoter driving expression of the non-native ORF6 can either be the native E4 promoter or a heterologous, non-native promoter operatively linked to the non-native ORF6, while in the latter, the expression of the non-native ORF6 would generally be driven by the native E4 promoter. Alternatively, the nucleic acid sequence encoding a partial E4 region comprising ORF 6 can be inserted into a different region of the adenoviral vector genome, such as for example the E1 or E3 regions. In this latter embodiment, the native E4 region or portion thereof can be deleted or left intact.

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As one of skill in the art can appreciate, there are various ways in which one can envision the supply of a heterologous E4 nucleic acid sequence *in cis* to an adenoviral vector and thereby enable its growth based on Applicants' novel findings herein. Moreover, as one of skill in the art can appreciate, either native or non-native promoters can be utilized to drive expression of the heterologous E4 region or portion thereof.

Adenovirus pre-plasmids (plasmids comprising the genome of the replication-defective adenovirus with desired deletions and insertions) can be generated by homologous recombination using adenovirus backbones and an appropriate shuttle vector (designed to targetin specific deletions and incorporate desired restriction sites into the resultant plasmid). Shuttle vectors of use in this process can be generated using general methods widely understood and appreciated in the art, e.g., PCR of the adenoviral terminal ends taking into account the desired deletions, and the sequential cloning of the respective segments into an appropriate cloning plasmid. The adenoviral pre-plasmid can then be digested and transfected into the complementing cell line via calcium phosphate co-precipitation or other suitable means. Virus replication and amplification then occurs, a phenomenon made evident by notable cytopathic effect. Infected cells and media are then harvested after viral replication is complete (generally, 7-10 days post-transfection).

It is to be noted that various alternative adenoviral serotypes can be developed in accordance with the disclosed methods and, particularly, alternative adenoviral serotype vectors that were previously unable to be propagated or very inefficiently propagated utilizing existing adenoviral production cell lines based on subgroup C complementing E1 sequence. The various adenoviral vectors that can be developed in accordance with the instant methods include adenoviral vectors of subgroups A-F (for instance, serotypes of subgroups A, B (e.g., serotypes

11, 14, 16, 21, 34 and 35), C (e.g., serotypes 2 and 5), D (e.g., serotypes 24, 26 and 36), E (e.g., serotype 4) and F.

In preferred embodiments, the various non-group C family members can be developed with heterologous E4 supplied from a subgroup C member such as adenovirus serotype 5. Particular embodiments of the instant invention utilize a development scheme wherein the heterologous E4 protein is derived from a wildtype adenovirus serotype 5 sequence; see, e.g., a viral sequence which has been deposited with the American Type Culture Collection ("ATCC") under ATCC Deposit No. VR-5 (for which a transcription map can be found in Figure 1). A particular example of this type of embodiment is wherein an adenovirus of subgroup B (or any non-C subgroup) comprising heterologous E4 proteins in cis from Ad5 is propagated in Ad5 E1-complementing cell lines, for instance, PER.C6TM or 293. Applicants have, in fact, successfully propagated E1- serotypes 10, 24, 34, and 35 via use of this particular embodiment.

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One of skill in the art can readily identify alternative adenovirus serotypes (e.g., alternative serotypes of subgroups A, B (e.g., serotypes 11, 14, 16, 21, 34 and 35), C, (e.g., serotypes 2 and 5), D (e.g., serotypes 24, 26 and 36), E (e.g., serotype 4) and F) for the supply of the heterologous E4 protein. As long as the heterologous E4 region (or portion thereof comprising ORF6) of the vector is native to a virus of the same or highly similar serotype as the E1 region of the complementing cell line, the methods of the instant invention are widely applicable to the propagation and rescue of adenovirus of all serotypes. In light of the present disclosure, one can readily envision, for instance, how a complementing cell line based on a non-subgroup C adenovirus (e.g., the Ad35 cell line of EP 1 054 064 A1) can be utilized to propagate a virus of an adenoviral vector of subgroup C (e.g., adenovirus serotype 5) provided that the appropriate nucleic acid sequence encoding an E4 protein provided in cis is native to a virus of the same or highly similar serotype as that of the E1 expressed by the complementing cell line (i.e., an Ad35 E4 protein).

Complementing cell lines of use in the instant invention are available in the art and are not limited to any specific type. The critical feature, again, is that the heterologous segment of E4-encoding nucleic acid sequence provided *in cis* to the replication-defective vector being propagated be native to a virus of the same or highly similar serotype as the E1 expressed by the complementing cell line. One aspect of the instant invention employs E1-complementing cell lines wherein the expressed E1 is of serotype 5; *e.g.*, PER.C6TM and 293 cell lines. Both these cell lines express the adenoviral E1 gene product. PER.C6TM is described in Fallaux *et al.*, 1998 *Human Gene Therapy* 9:1909-1917, hereby incorporated by reference. 293 cell lines are described in Graham *et al.*, 1977 *J. Gen. Virol.* 36:59-72, hereby incorporated by reference.

Another aspect of the instant invention are the adenoviral vectors of any serotype falling with adenoviral subgroups A, B, C, D, E and F (for instance, alternative serotypes of subgroups A, B (e.g., serotypes 11, 14, 16, 21, 34 and 35), C (e.g., serotype 2), D (e.g., serotypes 24, 26 and 36), E (e.g., serotype 4) and F) which are modified to contain a non-native E4-encoding nucleic acid sequence in cis which comprises, in whole or in part, nucleic acid sequence encoding open reading frame 6 (ORF6). Virus in accordance with this description can be propagated in accordance with the above-described methods and rescued using any suitable means known in the art.

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Another aspect of the instant invention is a vector in accordance with the instant invention which comprises a heterologous passenger gene in addition to that of the heterologous E4 nucleic acid sequence. In specific embodiments, the passenger gene encodes an antigen.

As one of ordinary skill in the art will appreciate, the instant methods are not limited by the heterologous gene that can be incorporated. The instant invention relates generally to a means by which to propagate multiple serotypes of adenovirus in a single complementing cell line and the recombinant virus that make the process possible. In preferred embodiments, the passenger gene is incorporated into the E1 deletion. In alternatively preferred embodiments, the passenger gene is inserted in an E3-deleted region. The position of the passenger gene, as one of ordinary skill in the art will appreciate, can be varied according to the specific complementing cell utilized and the specific deletions present within the replication-defective adenovirus genome.

In specific embodiments the passenger gene can encode an HIV-1 antigen, and in more preferred embodiments selected from the group consisting of genes encoding HIV-1 gag, pol, nef and env. In alternative embodiments, the passenger gene can be a reporter gene, such as secreted alkaline phosphatase (SEAP).

The passenger gene preferably exists in the form of an expression cassette. A gene expression cassette preferably comprises (a) a nucleic acid sequence encoding a protein of interest; (b) a promoter operatively linked to the nucleic acid sequence encoding the protein; and (c) a transcription termination sequence. The transcriptional promoter of the adenoviral vector is preferably recognized by an eukaryotic RNA polymerase. In a preferred embodiment, the promoter is a "strong" or "efficient" promoter. An example of a strong promoter is the immediate early human cytomegalovirus promoter (Chapman *et al.*, 1991 *Nucl. Acids Res.* 19:3979-3986), which is hereby incorporated by reference), in certain embodiments without intronic sequences. Those skilled in the art, however, will appreciate that any of a number of other known promoters, such as the strong immunoglobulin, or other eukaryotic gene promoters

may also be used, including the EF1 alpha promoter, the murine CMV promoter, Rous sarcoma virus (RSV) promoter, SV40 early/late promoters and the beta-actin promoter.

The promoter may comprise a regulatable sequence such as the Tet operator sequence. This is extremely useful, for example, in cases where the gene products are affecting a result other than that desired and repression is sought.

Transcription termination sequences can also be utilized within the gene expression cassettes. Preferred termination sequences are, for instance, the bovine growth hormone terminator/polyadenylation signal (bGHpA) and the short synthetic polyA signal (SPA) of 50 nucleotides in length, defined as follows:

10 NO:4).

Further embodiments incorporate a leader or signal peptide into the transgene. A preferred leader is that from the tissue-specific plasminogen activator protein, tPA.

The following non-limiting Examples are presented to better illustrate the

invention. 15

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EXAMPLE 1

Construction and Rescue

An E1- Ad35-based pre-adenovirus plasmid was constructed in order to determine whether an E1- Ad35 vector (a representative group B serotype) could be propagated in a group C E1-complementing cell line. The general strategy used to recover Ad35 as a bacterial plasmid is illustrated in Figure 3. Cotransformation of BJ5183 bacteria with purified wild-type Ad35 viral DNA and a second DNA fragment termed the Ad35 ITR cassette resulted in the circularization of the viral genome by homologous recombination. The ITR cassette contains sequences from the right (bp 34419 to 34793) and left (bp 4 to 456 and bp 3403 to 3886) end of the Ad35 genome (see Figures 2A-1 to 2A-10) separated by plasmid sequences containing a bacterial origin of replication and an Ampicillin resistance gene. The ITR cassette contains a deletion of E1 sequences from Ad5 457 to 3402 with a unique Swa I site located in the deletion. The Ad35 sequences in the ITR cassette provide regions of homology with the purified Ad35 viral DNA in which recombination can occur. The ITR cassette was also designed to contain unique restriction enzyme sites (Pme I) located at the end of the viral ITR's so that digestion will release the Ad35 genome from plasmid sequences. Potential clones were screened by restriction analysis and one clone was selected as pAd35ΔE1. Pre-Adenovirus plasmid pAd35ΔE1 contains Ad35 sequences from 4 to 456 and bp 3403 to 34793.

To determine if pre-adenovirus plasmid pAd35ΔE1 could be rescued into virus and propagated in a group C E1 complementing cell line, the plasmid was digested with *Pme* I and transfected into a T-25 flask of PER.C6 cells using the calcium phosphate co-precipitation technique. *Pme* I digestion releases the viral genome from the plasmid sequences allowing viral replication to occur after entry into 293 cells. Viral cytopathic effect (CPE), indicating that virus replication and amplification is occurring, was never observed. Cells and media from the transfection were harvested at 14 days post transfection, freeze-thawed three times, clarified by centrifugation and used to infect new PER.C6 cells but no virus was ever amplified. Following multiple attempts, we have been unable to rescue and amplify pAd35ΔE1 in PER.C6 cells.

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EXAMPLE 2

Insertion of Ad5 Orf 6 and Ad5 E4 into the Ad5 Genome

To refine the strategy of including Ad5 Orf6 in the genome of an alternative serotype so that propagation could take place in a Ad5/group C complementing cell line four additional strategies were developed. In the first strategy, the entire alternative serotype E4 region (not including the E4 promoter) was deleted and replaced with Ad5 Orf6. In the second strategy, just the alternative serotype Orf6 gene was deleted and replaced with Ad5 Orf6. In the third strategy, the entire alternative serotype E4 coding region (not including the E4 promoter) was deleted and replaced with the Ad5 E4 coding region (not including the Ad5 E4 promoter) and, in the final strategy, the entire alternative serotype E4 coding and promoter region was deleted and replaced with the Ad5 E4 promoter and coding region. The configuration of the E4 regions generated by the four strategies is diagramed in Figure 4. For each of these strategies the desired pre-Adenovirus plasmid was generated by bacterial recombination. Cotransformation of BJ 5183 bacteria with purified wild-type viral DNA and the appropriately constructed ITR cassette resulted in the circularization of the viral genome by homologous recombination. The construction of each pre-Ad plasmid, based on Ad35, is outlined below:

To construct pAd35 Δ E1 Δ E4Ad5Orf6 (An Ad35 pre-Ad plasmid containing an E1 deletion and an E4 deletion substituted with Ad5 Orf6), an Ad35 ITR cassette was constructed containing sequences from the right (bp 31599 to 31913 and bp 34419 to 34793) and left (bp 4 to 456 and bp 3403 to 3886) end of the Ad35 genome separated by plasmid sequences containing a bacterial origin of replication and an ampicillin resistance gene. These four segments were generated by PCR and cloned sequentially into pNEB193, generating pNEBAd35-4. Next the Ad5 Orf6 open reading frame was generated by PCR and cloned between Ad35 bp 31913 and 34419 generating pNEBAd35-4Ad5Orf6 (the ITR cassette). PNEB193 is a commonly used commercially available cloning plasmid (New England Biolabs cat# N3051S) containing a

bacterial origin of replication, ampicillin resistance gene and a multiple cloning site into which the PCR products were introduced. The ITR cassette contains a deletion of E1 sequences from Ad35 bp 457 to 3402 with a unique *Swa* I restriction site located in the deletion and an E4 deletion from Ad35 bp 31912 to 34418 into which Ad5 Orf6 was introduced in an E4 parallel orientation. In this construct, Ad5Orf6 expression is driven by the Ad35 E4 promoter. The Ad35 sequences (bp 31599 to 31913 and bp 3403 to 3886) in the ITR cassette provide regions of homology with the purified Ad35 viral DNA in which bacterial recombination can occur following cotransformation into BJ 5183 bacteria (Figure 5). The ITR cassette was also designed to contain unique restriction enzyme sites (PmeI) located at the end of the viral ITR's so that digestion will release the recombinant Ad35 genome from plasmid sequences. Potential clones were screened by restriction analysis and one clone was selected as pAd35ΔE1ΔE4Ad5Orf6. Pre-Adenovirus plasmid pAd35ΔE1ΔE4Ad5Orf6 contains Ad35 sequences from bp 4 to 456; bp 3403 to bp 31913 and bp 34419 to bp 34793 with Ad5Orf6 cloned between bp 31913 and bp 34419.

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To construct pAd35ΔE1ΔOrf6Ad5Orf6 (An Ad35 pre-Ad plasmid containing an E1 deletion and a deletion of E4 Orf6 substituted with Ad5 Orf6), an Ad35 ITR cassette was constructed containing sequences from the right (bp 31599 to 32081 and bp 32990 to 34793) and left (bp 4 to 456 and bp 3403 to 3886) end of the Ad35 genome separated by plasmid sequences containing a bacterial origin of replication and an ampicillin resistance gene. These four segments were generated by PCR and cloned sequentially into pNEB193, generating pNEBAd35-10. Next the Ad5 Orf6 open reading frame was generated by PCR and cloned between Ad35 bp 32081 and 32990 generating pNEBAd35-10Ad5Orf6 (the ITR cassette). PNEB193 is a commonly used commercially available cloning plasmid (New England Biolabs cat# N3051S) containing a bacterial origin of replication, ampicillin resistance gene and a multiple cloning site into which the PCR products were introduced. The ITR cassette contains a deletion of E1 sequences from Ad35 bp 457 to 3402 with a unique Swa I restriction site located in the deletion and a deletion of E4 Orf6 from Ad35 bp 32082 to 32989 into which Ad5 Orf6 was introduced in an E4 parallel orientation. In this construct, Ad5Orf6 expression is driven by the Ad35 E4 promoter. The Ad35 sequences (bp 31599 to 32081 and bp 3403 to 3886) in the ITR cassette provide regions of homology with the purified Ad35 viral DNA in which bacterial recombination can occur following cotransformation into BJ 5183 bacteria. The ITR cassette was also designed to contain unique restriction enzyme sites (Pme I) located at the end of the viral ITR's so that digestion will release the recombinant Ad35 genome from plasmid sequences. Potential clones were screened by restriction analysis and one clone was selected as pAd35ΔE1ΔOrf6Ad5Orf6. Pre-Adenovirus plasmid pAd35ΔE1ΔOrf6Ad5Orf6 contains Ad35

sequences from bp 4 to 456; bp 3403 to bp 32081 and bp 32990 to bp 34793 with Ad5Orf6 cloned between bp 32081 and bp 32990.

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To construct pAd35ΔE1ΔE4Ad5E4 (An Ad35 pre-Ad plasmid containing an E1 deletion and a deletion of E4 substituted with Ad5 E4), an Ad35 ITR cassette was constructed containing sequences from the right (bp 31599 to 31838 and bp 34419 to 34793) and left (bp 4 to 456 and bp 3403 to 3886) end of the Ad35 genome separated by plasmid sequences containing a bacterial origin of replication and an ampicillin resistance gene. These four segments were generated by PCR and cloned sequentially into pNEB193, generating pNEBAd35-7. Next the Ad5 E4 coding region was generated by PCR and cloned between Ad35 bp 31838 and 34419 generating pNEBAd35-7Ad5E4-2 (the ITR cassette). PNEB193 is a commonly used commercially available cloning plasmid (New England Biolabs cat# N3051S) containing a bacterial origin of replication, ampicillin resistance gene and a multiple cloning site into which the PCR products were introduced. The ITR cassette contains a deletion of E1 sequences from Ad35 bp 457 to 3402 with a unique Swa I restriction site located in the deletion and an E4 deletion from Ad35 bp 31839 to 34418 into which the Ad5 E4 coding region was introduced in an E4 parallel orientation. In this construct, the Ad5 E4 region is expressed using the Ad35 E4 promoter. The Ad35 sequences (bp 31599 to 31838 and bp 3403 to 3886) in the ITR cassette provide regions of homology with the purified Ad35 viral DNA in which bacterial recombination can occur following cotransformation into BJ 5183 bacteria. The ITR cassette was also designed to contain unique restriction enzyme sites (Pme I) located at the end of the viral ITR's so that digestion will release the recombinant Ad35 genome from plasmid sequences. Potential clones were screened by restriction analysis and one clone was selected as pAd35ΔE1ΔE4Ad5E4. Pre-Adenovirus plasmid pAd35ΔE1ΔE4Ad5E4 contains Ad35 sequences from bp 4 to 456; bp 3403 to bp 31838 and bp 34419 to bp 34793 with the Ad5 E4 coding region (Ad 5 bp 32914 to bp 35523) cloned between bp 31838 and bp 34419.

To construct pAd35ΔE1ΔE4Ad5PE4 (An Ad35 pre-Ad plasmid containing an E1 deletion and a deletion of E4 coding region and promoter substituted with Ad5 E4 coding region and promoter), an Ad35 ITR cassette was constructed containing sequences from the right (bp 31599 to 31838 and bp 34660 to 34793) and left (bp 4 to 456 and bp 3403 to 3886) end of the Ad35 genome separated by plasmid sequences containing a bacterial origin of replication and an ampicillin resistance gene. These four segments were generated by PCR and cloned sequentially into pNEB193, generating pNEBAd35-8. Next the Ad5 E4 promoter and coding region was generated by PCR and cloned between Ad35 bp 31838 and 34660 generating pNEBAd35-8Ad5E4PC (the ITR cassette). PNEB193 is a commonly used commercially available cloning plasmid (New England Biolabs cat# N3051S) containing a bacterial origin of replication,

ampicillin resistance gene, and a multiple cloning site into which the PCR products were introduced. The ITR cassette contains a deletion of E1 sequences from Ad35 bp 457 to 3402 with a unique Swa I restriction site located in the deletion and an E4 deletion from Ad35 bp 31839 to 34659 into which the Ad5 E4 promoter and coding region was introduced in an E4 parallel orientation. In this construct, the Ad5 E4 region is expressed using the Ad5 E4 promoter. The Ad35 sequences (bp 31599 to 31838 and bp 3403 to 3886) in the ITR cassette provide regions of homology with the purified Ad35 viral DNA in which bacterial recombination can occur following cotransformation into BJ 5183 bacteria. The ITR cassette was also designed to contain unique restriction enzyme sites (Pme I) located at the end of the viral ITR's so that digestion will release the recombinant Ad35 genome from plasmid sequences. Potential clones were screened by restriction analysis and one clone was selected as pAd35 Δ E1 Δ E4Ad5PE4. Pre-Adenovirus plasmid pAd35ΔE1ΔE4Ad5PE4 contains Ad35 sequences from bp 4 to 456; bp 3403 to bp 31838 and bp 34660 to bp 34793 with the Ad5 E4 promoter and coding region (Ad 5 bp 32914 to bp 35826) cloned between bp 31838 and bp 34660.

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EXAMPLE 3

Rescue of pAd35ΔE1ΔE4Ad5Orf6, pAd35ΔE1ΔOrf6Ad5Orf6, pAd35ΔE1ΔE4Ad5E4 and pAd35ΔE1ΔE4Ad5PE4 into Virus

In order to determine if pre-adenovirus plasmids pAd35ΔE1ΔE4Ad5Orf6, pAd35ΔE1ΔOrf6Ad5Orf6, pAd35ΔE1ΔE4Ad5E4 and pAd35ΔE1ΔE4Ad5PE4 could be rescued into virus and propagated in a group C E1 complementing cell line, the plasmids were each digested with Pme I and transfected into T-25 flasks of PER.C6 cells using the calcium phosphate co-precipitation technique; Cell Phect Transfection Kit, Amersham Pharmacia Biotech Inc. PmeI digestion releases the viral genome from plasmid sequences allowing viral replication to occur after cell entry. Viral cytopathic effect (CPE), indicating that virus replication and amplification was occurring, was observed for all construct. When CPE was complete, approximately 7-10 days post transfection, the infected cells and media were harvested, freeze/thawed three times and the cell debris pelleted by centrifugation. Approximately 1 ml of the cell lysate was used to infect aT-225 flasks of PER.C6 cells at 80-90% confluence. Once CPE was reached, infected cells and media were harvested, freeze/thawed three times and the cell debris pelleted by centrifugation. Clarified cell lysates were then used to infect 2-layer NUNC cell factories of PER.C6 cells. Following complete CPE the virus was purified by ultracentrifugation on CsCl density gradients. In order to verify the genetic structure of the rescued viruses, viral DNA was extracted using pronase treatment followed by phenol chloroform extraction and ethanol precipitation. Viral DNA was then

digested with *Hind*III and treated with Klenow fragment to end-label the restriction fragments with P33-dATP. The end-labeled restriction fragments were then size-fractionated by gel electrophoresis and visualized by autoradiography. The digestion products were compared with the digestion products of the corresponding pre-Adenovirus plasmid (that had been digested with *Pme1/Hind*III prior to labeling) from which they were derived. The expected sizes were observed, indicating that the viruses had been successfully rescued.

EXAMPLE 4

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Insertion of an Expression Cassette into pAd35ΔE1ΔE4Ad5Orf6, pAd35ΔE1ΔOrf6Ad5Orf6, pAd35ΔE1ΔE4Ad5E4 and pAd35ΔE1ΔE4Ad5PE4

In order to introduce a gag or SEAP expression cassette into the E1 region of the various Ad35 pre-Adenovirus plasmids described above (pAd35ΔE1ΔE4Ad5Orf6, pAd35ΔE1ΔOrf6Ad5Orf6, pAd35ΔE1ΔE4Ad5E4 and pAd35ΔE1ΔE4Ad5PE4) bacterial recombination was again used. A gag expression cassette consisting of the following: 1) the immediate early gene promoter from the human cytomegalovirus, 2) the coding sequence of the human immunodeficiency virus type 1 (HIV-1) gag (strain CAM-1; 1526 bp) gene, and 3) the bovine growth hormone polyadenylation signal sequence (Figure 6), was cloned into the E1 deletion in Ad35 shuttle plasmid, pNEBAd35-2 (a precursor to the Ad35 ITR cassettes described above), generating pNEBAd35CMVgagBGHpA. pNEBAd35-2 contains Ad35 sequences from the left end of the genome (bp 4 to 456 and bp 3403 to 3886) with a unique SwaI site between bp 456 and 3403 at the position of the deletion. The gag expression cassette was obtained from a previously constructed shuttle plasmid by EcoRI digestion. Following the digestion the desired fragment was gel purified, treated with Klenow to obtain blunt ends and cloned into the SwaI site in pNEBAd35-2. This cloning step resulted in the gag expression cassette being cloned into the E1 deletion between bp 456 and 3403 in the E1 parallel orientation. The shuttle vector containing the gag transgene was digested to generate a DNA fragment consisting of the gag expression cassette flanked by Ad35 bp 4 to 456 and bp 3403 to 3886 and the fragment was purified after electrophoresis on an agarose gel. Cotransformation of BJ 5183 bacteria with the shuttle vector fragment and one of the Ad35 pre-Ad plasmids (pAd35ΔE1ΔE4Ad5Orf6, pAd35ΔE1ΔOrf6Ad5Orf6, pAd35ΔE1ΔE4Ad5E4, pAd35ΔE1ΔE4Ad5PE4), linearized in the E1 region by digestion with Swa I, resulted in the generation of corresponding Ad35 gag-containing pre-Adenovirus plasmids (pAd35ΔE1gagΔE4Ad5Orf6, pAd35ΔE1gagΔOrf6Ad5Orf6, pAd35ΔE1gagΔE4Ad5E4, and pAd35ΔE1gagΔE4Ad5PE4) by homologous recombination.⁷ Potential clones were screened by restriction analysis.

A similar strategy was used to generate Ad35 pre-Ad plasmids containing a SEAP expression cassette. In this case a SEAP expression cassette consisting of: 1) the immediate early gene promoter from the human cytomegalovirus, 2) the coding sequence of the human placental SEAP gene, and 3) the bovine growth hormone polyadenylation signal sequence (Figure 7) was cloned into the E1 deletion in Ad35 shuttle plasmid, pNEBAd35-2, generating pNEBAd35CMVSEAPBGHpA. The SEAP expression cassette was obtained from a previously 5 constructed shuttle plasmid by EcoRI digestion. Following the digestion the desired fragment was gel purified, treated with Klenow to obtain blunt ends and cloned into the SwaI site in pNEBAd35-2. The transgene was then recombined into the various Ad35 backbones generating pAd35ΔE1SEAPΔE4Ad5Orf6, pAd35ΔE1SEAPΔOrf6Ad5Orf6, pAd35ΔE1SEAPΔE4Ad5E4, 10 and pAd35ΔE1SEAPΔE4Ad5PE4 as described above for the gag transgene. All pre-Ad plasmids were rescued into virus and expanded to prepare CsCl purified stocks as described above.

EXAMPLE 5 15 In vivo Transgene Expression

A. Immunization

Female mice were between 4-10 weeks old. The total dose of each vaccine was suspended in 0.1 mL of buffer. The vectors were given to both quadriceps of each animals with a volume of 50 μ L per quad and using 0.3-mL 28G1/2 insulin syringes (Becton-Dickinson, 20 Franklin Lakes, NJ). The rhesus macaques and African green monkeys were between 2-5 kg in weight. For the primates, the total dose of each vaccine was suspended in 1 mL of buffer. The monkeys were anesthetized (ketamine/xylazine mixture) and the vaccines were delivered i.m. in 0.5-mL aliquots into two muscle sites using tuberculin syringes (Becton-Dickinson, Franklin Lakes, NJ). Serum samples were collected at defined intervals and stored frozen until the assay 25 date. All animal care and treatment were in accordance with standards approved by the Institutional Animal Care and Use Committee according to the principles set forth in the Guide for Care and Use of Laboratory Animals, Institute of Laboratory Animal Resources, National Research Council. 30

B. SEAP Assay

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Serum samples were analyzed for circulating SEAP levels using TROPIX phospha-light chemiluminescent kit (Applied Biosystems Inc). Duplicate 5 µL aliquots of each serum were mixed with 45 μ L of kit-supplied dilution buffer in a 96-well white DYNEX plate.

Serially diluted solutions of a human placental alkaline phosphatase (Catalog no. M5905, Sigma, St. Louis, MO) in 10% naïve monkey or mouse serum served to provide the standard curve. Endogenous SEAP activity in the samples was inactivated by heating the well for 30 minutes at 65 °C. Enzymatic SEAP activities in the samples were determined following the procedures described in the kit. Chemiluminescence readings (in relative light units) were recorder using DYNEX luminometer. RLU readings are converted to ng/mL SEAP using a log-log regression analyses.

C. Rodent Results

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In the first mouse experiment, cohorts of 5 C3H/HeN mice were given single intramuscular injections of one of the following vectors: (1) 10^10 vp Ad35ΔE1SEAPΔE4Ad5Orf6; (2) 10^10 vp Ad35ΔE1SEAPΔE3ΔE4Ad5Orf6; or (3) 10^10 vp Ad35ΔE1SEAP. Serum samples prior to and after the injection were analyzed for circulating SEAP activities and the results are shown in Figure 8. Results indicate that (1) the Ad35 constructs are all capable of expressing the SEAP transgene and that (2) the introduction of Ad5Orf6 sequence where the deleted Ad35E4 was did not significantly affect the transgene expression relative to Ad35ΔE1SEAP. Ad35ΔE1SEAPΔE3ΔE4Ad5Orf6 also yielded a similar expression profile as Ad35ΔE1SEAP. The levels of SEAP in the serum dropped after day 2 and were at background levels by day 12.

The second mouse experiment evaluates the effect of a full Ad5E4 replacement instead of an Ad5Orf6 substitution for the Ad35 E4 cassette. Here, cohorts of 5 C3H/HeN mice were given single intramuscular injections of one of the following vectors: (1) 10^10 vp MRKAd5-SEAP; (2) 10^9 vp MRKAd5-SEAP; (3) 10^10 vp Ad35ΔE1SEAPΔE4Ad5Orf6; (4) 10^10 vp Ad35ΔE1SEAPΔE4Ad5E4; or (5) 10^10 vp Ad35ΔE1SEAPΔE4Ad5PE4. The introduction of Ad5E4 or Ad5PE4 resulted in comparable if not, slightly improved expression levels compared to the vector with the Ad5Orf6 insertion (Figure 9). The peak levels for the Ad35 constructs are lower than those produced by Ad5SEAP (at least 10-fold).

D. Primate Results

Cohorts of 3 rhesus macaques were given single intramuscular injections of one of the following vectors: (1) 10^11 vp MRKAd5-SEAP; (2) 10^9 vp MRKAd5-SEAP; or (3) 10^11 vp Ad35ΔE1SEAPΔE4Ad5Orf6. Serum samples prior to and after the injection were analyzed for circulating SEAP activities and the results for the individual monkeys are shown in Figures 10A-B. Results indicate that the peak level of SEAP product produced by the alternative adenovirus serotype was lower than but were within 3-fold of that of MRKAd5SEAP at the same

high dose level of 10^11 vp. The levels observed from the Ad35 vector were about 50-fold higher than those observed using 10^9 vp of MRKAd5SEAP. The levels of SEAP in the serum dropped after day 10 and were close to background as early as day 15.

A separate experiment using African green monkeys was conducted to examine the effect of the additional E3 deletion or the full Ad5E4 substitution on in vivo gene expression. In here, cohorts of 2-3 African green macaques were given single intramuscular injections of one 5 of the following vectors: (1) 10^11 vp MRKAd5-SEAP; (2) 10^10 vp MRKAd5-SEAP; (3) 10^9 vp MRKAd5-SEAP; (4) 10^10 vp Ad35ΔE1SEAPΔE4Ad5Orf6; (5) 10^10 vp Ad35ΔE1SEAPΔE3ΔE4Ad5Orf6; or (6) 10^10 vp Ad35ΔE1SEAPΔE4Ad5E4. Results (Figure 11) indicate that the peak levels of SEAP product produced by Ad35ΔE1SEAPΔE3ΔE4Ad5Orf6 and Ad35ΔE1SEAPΔE4Ad5E4 were comparable if not, slightly improved compared to 10 Ad35ΔE1SEAPΔE4Ad5Orf6.

EXAMPLE 6

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In vivo Immunogenicity 15

A. Immunization

Cohorts of 3-6 animals were given intramuscular injections at wk 0 and wk 4 of either of the following constructs: (1) 10^11 vp MRKAd5-HIV1 gag; or (2) 10^11 vp of Ad35ΔE1gagΔE4Ad5Orf6. Rhesus macaques were between 3-10 kg in weight. In all cases, the total dose of each vaccine was suspended in 1 mL of buffer. The macaques were anesthetized (ketamine/xylazine) and the vaccines were delivered i.m. in 0.5-mL aliquots into both deltoid muscles using tuberculin syringes (Becton-Dickinson). Sera and peripheral blood mononuclear cells (PBMC) were prepared from blood samples collected at several time points during the immunization regimen. All animal care and treatment were in accordance with standards approved by the Institutional Animal Care and Use Committee according to the principles set forth in the Guide for Care and Use of Laboratory Animals, Institute of Laboratory Animal Resources, National Research Council.

B. ELISPOT Assay

The IFN-γ ELISPOT assays for rhesus macaques were conducted following a previously described protocol (Allen et al., 2001 J. Virol. 75(2):738-749), with some modifications. For antigen-specific stimulation, a peptide pool was prepared from 20-aa peptides that encompass the entire HIV-1 gag sequence with 10-aa overlaps (Synpep Corp., Dublin, CA). To each well, 50 μ L of 2-4 x 10⁵ peripheral blood mononuclear cells (PBMCs)

were added; the cells were counted using Beckman Coulter Z2 particle analyzer with a lower size cut-off set at 80 femtoliters ("fL"). Either 50 μL of media or the gag peptide pool at 8 μg/mL concentration per peptide was added to the PBMC. The samples were incubated at 37°C, 5% CO₂ for 20-24 hrs. Spots were developed accordingly and the plates were processed using custom-built imager and automatic counting subroutine based on the ImagePro platform (Silver Spring, MD); the counts were normalized to 10⁶ cell input.

C. Intracellular Cytokine Staining

To 1 ml of 2 x 10⁶ PBMC/mL in complete RPMI media (in 17x100mm round bottom polypropylene tubes (Sarstedt, Newton, NC)), anti-hCD28 (clone L293, Becton-Dickinson) and anti-hCD49d (clone L25, Becton-Dickinson) monoclonal antibodies were added to a final concentration of 1 μ g/mL. For gag-specific stimulation, 10 μ L of the peptide pool (at 0.4 mg/mL per peptide) were added. The tubes were incubated at 37 °C for 1 hr., after which 20 μL of 5 mg/mL of brefeldin A (Sigma) were added. The cells were incubated for 16 hr at 37 °C, 5% CO₂, 90% humidity. 4 mL cold PBS/2%FBS were added to each tube and the cells were pelleted for 10 min at 1200 rpm. The cells were re-suspended in PBS/2%FBS and stained (30 min, 4 °C) for surface markers using several fluorescent-tagged mAbs: 20 μL per tube antihCD3-APC, clone FN-18 (Biosource); 20 μL anti-hCD8-PerCP, clone SK1 (Becton Dickinson, Franklin Lakes, NJ); and 20 μ L anti-hCD4-PE, clone SK3 (Becton Dickinson). Sample handling from this stage was conducted in the dark. The cells were washed and incubated in 750 μL 1xFACS Perm buffer (Becton Dickinson) for 10 min at room temperature. The cells were pelleted and re-suspended in PBS/2%FBS and 0.1 μg of FITC-anti-hIFN-γ, clone MD-1 (Biosource) was added. After 30 min incubation, the cells were washed and re-suspended in PBS. Samples were analyzed using all four color channels of the Becton Dickinson FACSCalibur instrument. To analyze the data, the low side- and forward-scatter lymphocyte population was initially gated; a common fluorescence cut-off for cytokine-positive events was used for both CD4⁺ and CD8⁺ populations, and for both mock and gag-peptide reaction tubes of a sample.

30 D. Results

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PBMCs collected at regular 4-wk intervals were analyzed in an ELISPOT assay. Results (Table 1) indicate that the Ad35ΔE1gagΔE4Ad5Orf6 is able to induce in non-human primates significant levels of gag-specific T cells. After a single dose (wk 4), the Ad35-induced responses were about 5-fold lower than that of MRKAd5-HIV1 gag. After the second dose (wk

8), the responses between both cohorts were comparable; the differences became pronounced in the succeeding time points.

Table 1. Gag-specific T cell response in monkeys immunized with MRKAd5-HIV1gag and Ad35ΔE1gagΔE4Ad5Orf6. Shown is the number of spot-forming cells per million PBMC following incubation in the absence (mock) or presence of Gag H peptide pool. The H pool consisted of 20-aa peptide overlapping by 10 aa and encompassing the entire gag sequence.

	Vaccine Wk 0, Wk 4	Monkey ID	Pre		Wk 4		Wk 8		Wk 12		Wk 16	
Gпр			Mock	Gag H	Mock	Gag H	Mock	Gag H	Mock	Gag H	Mock	Gag I
1	MRKAd5-HIV1 gag 10^11 vp	00C018 00C034 00C058	. 1 0 4	5 4 4	13 5 3	1025 219 1086	0 5 0	824 404 440	3 0 0	753 491 439	1 1 0	533 350 599
2	Ad35aE1gagAE4Ad5Orf6 10*11 vp	00D045 00D067 00D068 00D054 00D075 00D073	1 1 1 3 3 14	1 4 4 15 5 26	3 5 10 10 18 1	168 89 34 195 275 241	5 0 5 0 13 3	645 103 365 501 716 485	4 0 3 3 3 3	178 76 143 350 158 278	00000	91 19 95 124 103 148
3	Naïve	000087	1-1-	+-1	3	3	8	54	3	5	3	1_1

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Intracellular IFN-y staining analyses of PBMC collected at wk 8 suggest that the Ad35-based vaccine is able to induce both HIV-specific CD4+ and CD8+ T cells (Table 2).

Table 2. Characterization of the gag-specific T cells in monkeys immunized with MRKAd5-HIV1gag and Ad35∆E1gag∆E4Ad5Orf6. Shown are the percentages of CD3+ T cells that are 15 either gag-specific CD4+ or gag-specific CD8+ cells. These values were corrected for mock values (<0.02%).

	Vaccine	Monkey	Wk 8			
Grp	Wk 0, Wk 4	ID	%CD4+CD3+	%CD8+CD3+		
1	MRKAd5-HIV1 gag 10^11 vp	00C018 00C034 00C058	0.08 0.09 0.03	0.37 0.06 0.21		
2	Ad35∆E1gag∆E4Ad5Orf6 10^11 vp	00D045 00D067 00D068 00D054 00D075 00D073	0.06 0.02 0.15 0.05 0.08 0.09	0.08 0.02 0.02 0.08 0.05 0.06		

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In a separate experiment, 3 different Ad35 constructs expressing HIV-1 gag were evaluated for their immunogenicity in macaques. Here, cohorts of 3 macaques were given immunizations at wk 0 and 4 of either of the following vectors: (1) 10^10 vp Ad35∆E1gag∆E4Ad5Orf6; (2) 10^10

vp Ad35ΔE1gagΔE3ΔE4Ad5Orf6; or (3) 10^10 vp Ad35ΔE1gagΔE4Ad5E4. The levels of T cell immunity induced by all 3 vectors were comparable at this stage (Table 2), suggesting that the additional E3 deletion or full Ad5E4 substitution does not appear to impair the immunogenic properties of the vector.

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Table 3. Gag-specific T cell response in monkeys immunized with several Ad35ΔE1ΔE4-based vectors. Shown is the number of spot-forming cells per million PBMC following incubation in the absence (mocK0 or presence of Gag H peptide pool. The H pool consisted of 20-aa peptide overlapping by 10 aa and encompassing the entire gag sequence.

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Grp	Vaccine	Monkey	Pre		Wk 4		Wk 8	
Gib	Wk 0, Wk 4	ID	Mock	Gag H	Mock	Gag H	Mock	Gag H
	Ad35∆E1gag∆E4Ad5Orf6	00C047	4	1	0	20	0	189
'	10^10vp	00C157	8	5	1	81	1	833
	10 104μ	00C078	3	1	0	46	4	349
2	Ad35∆E1gag∆E3∆E4Ad5Orf6	00C091	1	1	1	118	3	315
- 1	10^10vp	00C122	3	0	0	31	1	138
	10 1000	00D177	3	3	1	45	1	64
	Ad35∆E1gag∆E4Ad5E4	00D018	3	19	29	120	23	193
3	10*10vp	00D046	8	5	1 1	21	10	143
	ιστιούρ	00D063	3	4	0	63	4	371
		<u> </u>			- 115	NID	0	0
Naîve	none	00D363	0	5	ND	ND	<u> </u>	

EXAMPLE 7

Construction and Rescue of pAd24ΔE1.

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An E1- Ad24-based pre-adenovirus plasmid was constructed in order to determine whether an E1- Ad24 vector (a representative group D serotype) could be propagated in an Ad5/group C E1-complementing cell line. Since at the time the vector construction was initiated the complete sequence of Ad24 (see Figures 16A-1 through 16A-10; subject of copending application serial no. 60/455, 312, filed March 17, 2003) was unknown we took advantage of some sequence homology between Ad24 and Ad17. The general strategy used to recover Ad24 as a bacterial plasmid is illustrated in Figure 12 and described below. Cotransformation of BJ5183 bacteria with purified wild-type Ad24 viral DNA and a second DNA fragment termed the Ad17 ITR cassette resulted in the circularization of the viral genome by homologous recombination. The ITR cassette contains sequences from the right (bp 34469 to 35098) and left (bp 4 to 414 and bp 3373 to 4580) end of the Ad17 genome (Accession No. AF108105) separated by plasmid sequences containing a bacterial origin of replication and an Ampicillin resistance gene. The ITR cassette contains a deletion of E1 sequences from Ad17

(bp 415 to 3372) with a unique Swa I site located in the deletion. The Ad17 sequences in the ITR cassette provide regions of homology with the purified Ad24 viral DNA in which recombination can occur. The ITR cassette was also designed to contain unique restriction enzyme sites (Pme I) located at the end of the viral ITR's so that digestion will release the Ad24 genome from plasmid sequences. Potential clones were screened by restriction analysis and one clone was selected as pAd24ΔE1. pAd24ΔE1 contains Ad17 sequences from bp 4 to 414 and from bp 3373 to 4580, Ad24 bp 4588 to 34529, and Ad17 bp 34469 to 35098 (bp numbers refer to the wt sequence for both Ad17 and Ad24). PAd24ΔE1 contains the coding sequences for all Ad24 virion structural proteins that constitute its serotype specificity. This approach can be used to circularize any group D serotype into plasmid form which has sufficient homology to Ad17.

To determine if pre-adenovirus plasmid pAd24ΔE1 could be rescued into virus and propagated in a group C E1 complementing cell line, the plasmid was digested with *Pme* I and transfected into a 6 cm dish of 293 cells using the calcium phosphate co-precipitation technique. *Pme* I digestion releases the viral genome from the plasmid sequences allowing viral replication to occur after entry into 293 cells. Viral cytopathic effect (CPE), indicating that virus replication and amplification is occurring, was very slow to arise. Following multiple attempts, we were successful at rescuing and amplifying Ad24ΔE1 but the virus grew to lower titers and took more passages to amplify than a similar Ad5 based vector. In order to verify the genetic structure of the virus, viral DNA was extracted using pronase treatment followed by phenol chloroform extraction and ethanol precipitation. Viral DNA was then digested with *Hind*III and treated with Klenow fragment to end-label the restriction fragments with P33-dATP. The end-labeled restriction fragments were then size-fractionated by gel electrophoresis and visualized by autoradiography. The digestion products were compared with the digestion products from the pre-plasmid (that had been digested with *Pme1/Hind*III prior to labeling). The expected sizes were observed, indicating that the virus had been successfully rescued.

EXAMPLE 8

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Insertion of Ad5 Orf 6 into the E1 region of Ad24

In order to determine if the insertion of Ad5 E4 Orf6 into the Ad24 genome would allow more efficient propagation in a group C E1 complementing cell line we constructed an Ad24 based pre-adenovirus plasmid containing Ad5 Orf6 in the E1 region. In order to introduce Ad5 Orf6 in to the E1 region of pAd24ΔE1, bacterial recombination was used. An Ad5 Orf6 transgene consisting of the Ad5 Orf6 coding region flanked by the HCMV promoter and pA was cloned into the E1 deletion in an Ad17 shuttle vector (a precursor to the Ad17 ITR cassette). The Ad5 Orf6 transgene was cloned between bp 414 and 3373 in the E1 anti-parallel

orientation. The shuttle vector containing the Ad5 Orf6 transgene was digested to generate a DNA fragment consisting of the transgene flanked by Ad17 sequences (bp 4 to 414 and bp 3373 to 4580) and the fragment was purified after electrophoresis on an agarose gel. Cotransformation of BJ 5183 bacteria with the shuttle vector fragment and pAd24ΔE1, which had been linearized in the E1 region by digestion with *Swa*I, resulted in the generation of pAd24ΔE1Ad5Orf6 by homologous recombination (Figure 13). Potential clones were screened by restriction analysis and one clone was selected as pre-adenovirus plasmid pAd24ΔE1Ad5Orf6.

In order to determine if pre-adenovirus plasmid pAd24ΔE1Ad5Orf6 could be rescued into virus and propagated in an Ad5/group C E1 complementing cell line, pAd24ΔE1Ad5Orf6 was digested with Pme I and transfected into a 6 cm dish of 293 cells using the calcium phosphate co-precipitation technique. PmeI digestion releases the viral genome from plasmid sequences allowing viral replication to occur after entry into 293 cells. Once complete viral cytopathic effect (CPE) was observed at approximately 7-10 days post transfection, the infected cells and media were freeze/thawed three times and the cell debris pelleted. The virus was amplified in two additional passages in 293 cells and then purified from the final infection by ultracentrifugation on CsCl density gradients. In order to verify the genetic structure of the virus, viral DNA was extracted using pronase treatment followed by phenol chloroform extraction and ethanol precipitation. Viral DNA was then digested with HindIII and treated with Klenow fragment to end-label the restriction fragments with P33-dATP. The endlabeled restriction fragments were then size-fractionated by gel electrophoresis and visualized by autoradiography. The digestion products were compared with the digestion products from the pre-plasmid (that had been digested with Pme1/HindIII prior to labeling). The expected sizes were observed, indicating that the virus had been successfully rescued.

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EXAMPLE 9

Insertion of Ad5 Orf 6 into the E4 region of Ad24

To refine the strategy of including Ad5 Orf6 in the genome of an alternative serotype so that propagation could take place in an Ad5/group C complementing cell line two additional strategies were developed. In the first strategy, the entire alternative serotype E4 region (not including the E4 promoter) was deleted and replaced with Ad5 Orf6. In the second strategy, just the alternative serotype Orf6 gene was deleted and replaced with Ad5 Orf6. The configuration of the E4 regions generated by the two strategies is diagramed in Figure 14. For each of these strategies the desired pre-Adenovirus plasmid was generated by bacterial recombination. Cotransformation of BJ 5183 bacteria with pAd24ΔOrf6BstZ17I and the

appropriately constructed Ad24 E4 shuttle plasmid resulted in the generation of the desired Ad24 based pre-Ad plasmid. PAd24ΔOrf6BstZ17I, a derivative of pAd24ΔE1, was constructed so that the E4 region in the Ad24 pre-Ad plasmid could be easily modified using bacterial recombination. PAd24ΔOrf6BstZ17I contains a deletion in the E4 region from Ad24 bp 32373 to bp 33328 with a unique BstZ17I site located at the position of the deletion. The complete sequence of pAd24ΔOrf6BstZ17I consists of Ad17 sequences from bp 4 to 414 and from bp 3373 to 4580, Ad24 bp 4588 to 32372 and from 33329 to 34529, and Ad17 bp 34469 to 35098 (bp numbers refer to the wt sequence for both Ad17 and Ad24).

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To construct pAd24ΔE1ΔE4Ad5Orf6 (An Ad24 pre-Ad plasmid containing an E1 deletion and a deletion of E4 substituted with Ad5 Orf6), an Ad24 E4 shuttle plasmid was constructed by digesting pAd24ΔE1 with *PmeI* and *BsrGI* and cloning the restriction fragment representing the E4 region (bp 31559 to bp 35164) into pNEB193, generating pNEBAd24E4. PNEBAd24E4 was then digested with *AccI* and *Eco*NI to remove the E4 coding sequences and ligated with an oligo designed to contain *BgIII* and *XhoI* sites (underlined) (5'

- 15 ACTCGAGATGTATAGATCT (SEQ ID NO: 6); 5' CTAGATCTATACATCTCGAG (SEQ ID NO: 7)), generating pNEBAd24ΔE4. PNEBAd24ΔE4 was then digested with Bg/II and XhoI and ligated with the Ad5 Orf6 gene, which was PCR amplified, generating pNEBAd24ΔE4Ad5Orf6. The PCR primers used to amplify the Ad5 Orf6 gene (5' GCACAGATCTTTGCTTCAGGAATATG (SEQ ID NO: 8); 5'
- GAGAACTCGAGGCCTACATGGGGGTAGAG (SEQ ID NO: 9)) were designed to contain BgIII and XhoI sites (underlined above) for ligation with the pNEBAd24DE4 fragment. In the final step pNEBAd24ΔE4Ad5Orf6 E4 shuttle plasmid was digested with PvuI and PmeI, the restriction fragments were size fractionated by agarose gel electrophoresis and the desired fragment containing Ad5Orf6 flanked by Ad24 sequences was gel purified. Cotransformation of BJ 5183 bacteria with E4 shuttle fragment and pAd24ΔOrf6BstZ17I, which had been linearized in the E4 region by digestion with BstZ17I, resulted in the generation of pAd24ΔE1ΔE4Ad5Orf6 by homologous recombination. Potential clones were screened by restriction analysis and one clone was selected as pre-adenovirus plasmid pAd24ΔE1ΔE4Ad5Orf6.

To construct pAd24ΔE1ΔOrf6Ad5Orf6 (An Ad24 pre-Ad plasmid containing an E1 deletion and a deletion of E4 Orf6 substituted with Ad5 Orf6), an Ad24 E4 shuttle plasmid was constructed in which the Ad24 Orf6 gene was replaced by Ad5 Orf6. To do this the *Eco*R1 restriction fragment representing bp 32126 to bp 33442 of the Ad24 genome (encompassing the E4 Orf6 coding region), was subcloned into the *Eco*RI site in pNEB193, generating pNEBAd24Orf6. In order to delete the E4 Orf6 gene in pNEBAd24Orf6 and replace it with Ad5 Orf6, pNEBAd24Orf6 was digested with *Sty*I and treated with Klenow to blunt the ends and then

digested with to EagI. The desired pNEBAd24Orf6 fragment was then ligated with a PCR product representing the Ad5 Orf6 gene from Ad5 bp 33193 to bp 24125, generating pNEBAd24ΔOrf6Ad5Orf6. The PCR primers used to generate the Ad5 Orf6 fragment (5'CGAGACGCCGACGCAGATCTGTTTG (SEQ ID NO: 10);

5'GAAGTCCCGGGCTACATGGGGGTAG (SEQ ID NO: 11)) were designed to contain EagI and SmaI sites (underlined above) for ligation with the pNEBAd24Orf6 fragment. In the final step pNEBAd24ΔOrf6Ad5Orf6 was digested with EcoRI, the restriction fragments were size fractionated by agarose gel electrophoresis and the desired fragment containing Ad5Orf6 flanked by Ad24 sequences was gel purified. Cotransformation of BJ 5183 bacteria with the EcoRI fragment and pAd24ΔOrf6BstZ17I, which had been linearized in the E4 region by digestion with BstZ17I, resulted in the generation of pAd24ΔE1ΔOrf6Ad5Orf6 by homologous recombination. Potential clones were screened by restriction analysis and one clone was selected as preadenovirus plasmid pAd24ΔE1ΔOrf6Ad5Orf6.

15 EXAMPLE 10 Rescue of pAd24ΔE1ΔE4Ad5Orf6, pAd24ΔE1ΔOrf6Ad5Orf6, into Virus

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pAd24 Δ E1 Δ Orf6Ad5Orf6, could be rescued into virus and propagated in a group C E1 complementing cell line, the plasmids were each digested with *Pme* I and transfected into T-25 flasks of PER.C6 cells using the calcium phosphate co-precipitation technique; (Cell Phect Transfection Kit, Amersham Pharmacia Biotech Inc.). *PmeI* digestion releases the viral genome from plasmid sequences allowing viral replication to occur after cell entry. Viral cytopathic effect (CPE), indicating that virus replication and amplification was occurring, was observed for both constructs. When CPE was complete, approximately 7-10 days post transfection, the infected cells and media were harvested, freeze/thawed three times and the cell debris pelleted by centrifugation. Approximately 1 ml of the cell lysate was used to infect T-225 flasks of PER.C6 cells at 80-90% confluence. Once CPE was reached, infected cells and media were

In order to determine if pre-adenovirus plasmids pAd24\Delta E1\Delta E4Ad5Orf6,

complete CPE the virus was purified by ultracentrifugation on CsCl density gradients. In order to verify the genetic structure of the rescued viruses, viral DNA was extracted using pronase treatment followed by phenol chloroform extraction and ethanol precipitation. Viral DNA was then digested with *Hin*dIII and treated with Klenow fragment to end-label the restriction fragments with P33-dATP. The end-labeled restriction fragments were then size-fractionated by gel electrophoresis and visualized by autoradiography. The digestion products were compared

harvested, freeze/thawed three times and the cell debris pelleted by centrifugation. Clarified cell

lysates were then used to infect 2-layer NUNC cell factories of PER.C6 cells. Following

with the digestion products of the corresponding pre-Adenovirus plasmid (that had been digested with Pme1/HindIII prior to labeling) from which they were derived. The expected sizes were observed, indicating that the viruses had been successfully rescued.

EXAMPLE 11 5

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Comparison of the Growth Kinetics of Ad24 based vectors.

In order to compare the growth kinetic of Ad24 \Delta E1, Ad24 \Delta E1 Ad5 Orf6, Ad24ΔE1ΔE4Ad5Orf6 and Ad24ΔE1ΔOrf6Ad5Orf6 one step growth curves were preformed (Figure 15). PER.C6 cells in 60 mm dishes were infected at 1 vp per cell with either Ad24ΔE1, Ad24ΔE1Ad5Orf6, Ad24ΔE1ΔE4Ad5Orf6 or Ad24ΔE1ΔOrf6Ad5Orf6. Cells and media were then harvested at various times post infection, freeze thawed three times and clarified by centrifugation. The amount of virus present in the samples was determined by quantitative PCR and is illustrated in Figure 15. This study demonstrates that Ad24 vectors that incorporate Ad5 Orf6 have a distinct growth advantage over Ad24 Δ E1 in PER.C6 cells. The instant invention can be practiced with recombinant Ad24 vectors absent a heterologous Orf 6 region where the E1-complementing cell line expresses an Ad24 E1 region or, alternatively, E1 and E4 regions of the same serotype (such as Ad5E1/E4-expressing cell lines).

EXAMPLE 12

Insertion of an Expression Cassette into pAd24ΔE1ΔE4Ad5Orf6, pAd24ΔE1ΔOrf6Ad5Orf6, In order to introduce a gag or SEAP expression cassette (see Figures 6 and 7, 20 respectively) into the E1 region of the Ad24 pre-Adenovirus plasmids described above (pAd24ΔE1ΔE4Ad5Orf6, pAd24ΔE1ΔOrf6Ad5Orf6) bacterial recombination was used. A gag expression cassette consisting of the following: 1) the immediate early gene promoter from the human cytomegalovirus, 2) the coding sequence of the human immunodeficiency virus type 1 (HIV-1) gag (strain CAM-1; 1526 bp) gene, and 3) the bovine growth hormone polyadenylation 25 signal sequence, was cloned into the E1 deletion in Ad17 shuttle plasmid, pABSAd17-3, generating pABSAd17HCMVgagBGHpA. The ITR cassette contains sequences from the right (bp 34469 to 35098) and left (bp 4 to 414 and bp 3373 to 4580) end of the Ad17 genome separated by plasmid sequences containing a bacterial origin of replication and an Ampicillin resistance gene. The ITR cassette contains a deletion of E1 sequences from Ad17 (bp 415 to 30 3372) with a unique Swa I site located in the deletion. The gag expression cassette was obtained from a previously constructed shuttle plasmid by EcoRI digestion. Following the digestion the desired fragment was gel purified, treated with Klenow to obtain blunt ends and cloned into the Swal site in pABSAd17-3. This cloning step resulted in the gag expression cassette being 35

cloned into the E1 deletion between bp 414 and 3373 in the E1 parallel orientation. The shuttle vector containing the gag transgene was digested to generate a DNA fragment consisting of the gag expression cassette flanked by Ad17 bp 4 to 414 and bp 3373 to 4580 and the fragment was purified after electrophoresis on an agarose gel. Cotransformation of BJ 5183 bacteria with the shuttle vector fragment and one of the Ad24 pre-Ad plasmids (pAd24ΔE1ΔE4Ad5Orf6, pAd24ΔE1ΔOrf6Ad5Orf6,), linearized in the E1 region by digestion with Swa I, resulted in the generation of the corresponding Ad24 gag-containing pre-Adenovirus plasmids (pAd24ΔE1gagΔE4Ad5Orf6, pAd24ΔE1gagΔOrf6Ad5Orf6) by homologous recombination. Potential clones were screened by restriction analysis.

A similar strategy was used to generate Ad24 pre-Ad plasmids containing a SEAP expression cassette. In this case a SEAP expression cassette consisting of: 1) the immediate early gene promoter from the human cytomegalovirus, 2) the coding sequence of the human placental SEAP gene, and 3) the bovine growth hormone polyadenylation signal sequence was cloned into the E1 deletion in Ad17 shuttle plasmid, pABSAd17-3, generating pABSAd17HCMVSEAPBGH. The SEAP expression cassette was obtained from a previously constructed shuttle plasmid by EcoRI digestion. Following the digestion the desired fragment was gel purified, treated with Klenow to obtain blunt ends and cloned into the SwaI site in pABSAd17-3. The shuttle vector containing the SEAP transgene was digested to generate a DNA fragment consisting of the SEAP expression cassette flanked by Ad17 bp 4 to 414 and bp 3373 to 4580 and the fragment was purified after electrophoresis on an agarose gel. Cotransformation of BJ 5183 bacteria with the shuttle vector fragment and one of the Ad24 pre-Ad plasmids (pAd24ΔE1ΔE4Ad5Orf6, pAd24ΔE1ΔOrf6Ad5Orf6,), linearized in the E1 region by digestion with Swa I, resulted in the generation of the corresponding Ad24 SEAP-containing pre-Adenovirus plasmids (pAd24ΔE1SEAPΔE4Ad5Orf6, pAd24ΔE1SEAPΔOrf6Ad5Orf6) by homologous recombination. Potential clones were screened by restriction analysis. All pre-Ad plasmids were rescued into virus and expanded to prepare CsCl purified stocks as described above.

EXAMPLE 13

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30 In Vivo Immunogenicity

A. Immunization

Cohorts of 3-6 animals were given intramuscular injections at wk 0 and wk 4 of either of the following constructs: (1) 10^11 vp MRKAd5-HIV1 gag; (2) 10^10 vp MRKAd5-HIV1 gag; (3) 10^11 vp of Ad24ΔE1gagΔOrf6Ad5Orf6; (4) 10^10 vp of

Ad24ΔE1gagΔOrf6Ad5Orf6; or (5) 10^10 vp of Ad24ΔE1gagΔE4Ad5Orf6. Rhesus macaques were between 3-10 kg in weight. In all cases, the total dose of each vaccine was suspended in 1 mL of buffer. The macaques were anesthetized (ketamine/xylazine) and the vaccines were delivered i.m. in 0.5-mL aliquots into both deltoid muscles using tuberculin syringes (Becton-Dickinson, Franklin Lakes, NJ). Peripheral blood mononuclear cells (PBMC) were prepared from blood samples collected at several time points (typically 4 wk intervals) during the immunization regimen. All animal care and treatment were in accordance with standards approved by the Institutional Animal Care and Use Committee according to the principles set forth in the Guide for Care and Use of Laboratory Animals, Institute of Laboratory Animal Resources, National Research Council.

B. ELISPOT Assay

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The IFN-γ ELISPOT assays for rhesus macaques were conducted following a previously described protocol (Allen et al., 2001 J. Virol. 75(2):738-749; Casimiro et al., 2002 J. Virol. 76:185-94), with some modifications. For antigen-specific stimulation, a peptide pool was prepared from 20-aa peptides that encompass the entire HIV-1 gag sequence with 10-aa overlaps (Synpep Corp., Dublin, CA). To each well, 50 μL of 2-4 x 10⁵ peripheral blood mononuclear cells (PBMCs) were added; the cells were counted using Beckman Coulter Z2 particle analyzer with a lower size cut-off set at 80 femtoliters ("fL"). Either 50 µL of media or the gag peptide pool at 8 µg/mL concentration per peptide was added to the PBMC. The samples were incubated at 37°C, 5% CO₂ for 20-24 hrs. Spots were developed accordingly and the plates were processed using custom-built imager and automatic counting subroutine based on the ImagePro platform (Silver Spring, MD); the counts were normalized to 10^6 cell input.

C. Intracellular Cytokine Staining 25

To 1 ml of 2 x 10^6 PBMC/mL in complete RPMI media (in 17x100mm round bottom polypropylene tubes (Sarstedt, Newton, NC)), anti-hCD28 (clone L293, Becton-Dickinson) and anti-hCD49d (clone L25, Becton-Dickinson) monoclonal antibodies were added to a final concentration of 1 μ g/mL. For gag-specific stimulation, 10 μ L of the peptide pool (at 0.4 mg/mL per peptide) were added. The tubes were incubated at 37 °C for 1 hr., after which 20 μL of 5 mg/mL of brefeldin A (Sigma) were added. The cells were incubated for 16 hr at 37 °C, 5% CO₂, 90% humidity. 4 mL cold PBS/2%FBS were added to each tube and the cells were pelleted for 10 min at 1200 rpm. The cells were re-suspended in PBS/2%FBS and stained (30 min, 4 °C) for surface markers using several fluorescent-tagged mAbs: 20 μL per tube antihCD3-APC, clone FN-18 (Biosource); 20 μL anti-hCD8-PerCP, clone SK1 (Becton Dickinson);

and 20 μ L anti-hCD4-PE, clone SK3 (Becton Dickinson). Sample handling from this stage was conducted in the dark. The cells were washed and incubated in 750 μ L 1xFACS Perm buffer (Becton Dickinson) for 10 min at room temperature. The cells were pelleted and re-suspended in PBS/2%FBS and 0.1 μ g of FITC-anti-hIFN- γ , clone MD-1 (Biosource) was added. After 30 min incubation, the cells were washed and re-suspended in PBS. Samples were analyzed using all four color channels of the Becton Dickinson FACSCalibur instrument. To analyze the data, the low side- and forward-scatter lymphocyte population was initially gated; a common fluorescence cut-off for cytokine-positive events was used for both CD4⁺ and CD8⁺ populations, and for both mock and gag-peptide reaction tubes of a sample.

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D. Anti-p24 ELISA

A modified competitive anti-p24 assay was developed using reagents from the Coulter p24 Antigen Assay kit (Beckman Coulter, Fullerton, CA). Briefly, to a 250-μL serum sample, 20 μL of Lyse Buffer and 15 μL of p24 antigen (9.375 pg) from the Coulter kit were added. After mixing, 200 μL of each sample were added to wells coated with a mouse anti-p24 mAb from the Coulter kit and incubated for 1.5 hr at 37°C. The wells were then washed and 200 μL of Biotin Reagent (polyclonal anti-p24-biotin) from the Coulter kit was added to each well. After a 1 hr, 37°C incubation, detection was achieved using strepavidin-conjugated horseradish peroxidase and TMB substrate as described in the Coulter Kit. OD450nm values were recorded. A 7-point standard curve was generated using a serial 2-fold dilution of serum from an HIV-seropositive individual. The lower cut-off for the assay is arbitrarily set at 10 milli Merck units/mL (mMU/mL) defined by a dilution of the seropositive human serum. This cutoff falls at approximately 65% of the maximum bound control signal which corresponds to that obtained with the diluent control only and with no positive analyte.

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E. Results

PBMCs collected at regular 4-wk intervals were analyzed in an ELISPOT assay (Figure 17). Both Ad24ΔE1gagΔOrf6Ad5Orf6 and Ad24ΔE1gagΔE4Ad5Orf6 were able to induce significant levels of gag-specific T cells in non-human primates. At 10^11 vp dose level, the Ad24-induced responses were within 2-3-fold of those of MRKAd5-HIV1 gag. Both Ad24 vectors were also able to induce detectable levels of gag-specific T cells at 10^10 vp but were lower than those observed using MRKad5gag at the same dose.

PBMCs collected at wk 12 from the vaccinees were analyzed for intracellular IFN-γ staining after the priming immunizations. The assay results provided information on the relative amounts of CD4⁺ and CD8⁺ gag-specific T cells in the peripheral blood (Figure 18). The

results indicated that the prime-boost immunization approach was able to elicit in rhesus macaques both HIV-specific CD4⁺ and CD8⁺ T cells.

F. Humoral Immune Responses

The Ad24-based vaccine vector was able to generate detectable levels of circulating anti-gag antibodies at the reasonably high dose level (Figure 19). No detectable titers were observed at equal to or lower than 10^10 vp, suggesting the existence of a dose-dependent response.

EXAMPLE 14 10

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In Vivo Transgene Expression

A. Immunization

Cohorts of 5 C3H/HeN mice were given single intramuscular injections of one of the following vectors: (1) 10^10 vp Ad24 Δ E1SEAP Δ E4Ad5Orf6; (2) 10^10 vp Ad24ΔE1SEAPΔOrf6Ad5Orf6; (3) 10^10 vp MRKAd5SEAP; and (4) 10^9 vp MRKAd5SEAP. Female mice were between 4-10 weeks old. The total dose of each vaccine was suspended in 0.1 mL of buffer. The vectors were given to both quadriceps of each of the animals with a volume of 50 uL per quad and using 0.3-mL 28G1/2 insulin syringes (Becton-Dickinson, Franklin Lakes, NJ). For the primates, the total dose of each vaccine was suspended in 1 mL of buffer. The monkeys were anesthetized (ketamine/xylazine mixture) and the vaccines were delivered i.m. in 20 0.5-mL aliquots into two muscle sites using tuberculin syringes (Becton-Dickinson, Franklin Lakes, NJ). Serum samples were collected at defined intervals and stored frozen until the assay date. All animal care and treatment were in accordance with standards approved by the Institutional Animal Care and Use Committee according to the principles set forth in the Guide for Care and Use of Laboratory Animals, Institute of Laboratory Animal Resources, National 25 Research Council.

B. SEAP Assay

Serum samples were analyzed for circulating SEAP levels using TROPIX phospha-light chemiluminescent kit (Applied Biosystems Inc). Duplicate 5 uL aliquots of each serum were mixed with 45 uL of kit-supplied dilution buffer in a 96-well white DYNEX plate. Serially diluted solutions of a human placental alkaline phosphatase (Catalog no. M5905, Sigma, St. Louis, MO) in 10% naïve monkey serum served to provide the standard curve. Endogenous SEAP activity in the samples was inactivated by heating the wells for 30 minutes at 65 °C.

Enzymatic SEAP activities in the samples were determined following the procedures described in the kit. Chemiluminescence readings (in relative light units) were recorder using DYNEX luminometer. RLU readings are converted to ng/mL SEAP using a log-log regression analyses.

5 C. Rodent Results

Serum samples prior to and after the injection were analyzed for circulating SEAP activities and the results are shown in Figure 20. Results indicate that (1) both Ad24 constructs are all capable of expressing the SEAP transgene in vivo to comparable levels; and that (2) the level of expression achieved using the Ad24 vectors are comparable to that of Ad5 at 10-fold lower dose. The levels of SEAP in the serum dropped dramatically after day 2 and were at background levels by day 12.

D. Primate Results

Cohorts of 3 rhesus macaques were given single intramuscular injections of one of the following vectors: (1) 10^11 vp MRKAd5-SEAP; (2) 10^9 vp MRKAd5-SEAP; (3) 10^11 vp Ad24ΔE1SEAPΔOrf6Ad5Orf6; or (4) 10^11 vp Ad24ΔE1SEAPΔE4Ad5Orf6. Serum samples prior to and after the injection were analyzed for circulating SEAP activities and the results are shown in Figure 21.

Results indicate that the peak levels of SEAP product produced by adenovirus serotype 24 were lower than but were within 3-fold of that of MRKAd5 at the same high dose level of 10^11 vp (Figure 21). The levels observed with adenovirus serotype 24 are generally 50-fold higher than those observed using 10^9 vp of MRKAd5. The levels of SEAP in the serum dropped dramatically after day 10 and were close to background as early as day 15. These observations strongly indicate that adenovirus serotype 24 is very efficient in expressing a transgene following intramuscular administration in a primate.

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EXAMPLE 15

Construction of pMRKAd24ΔE1ΔE4Ad5Orf6

To construct pMRKAd24ΔE1ΔE4Ad5Orf6 (An Ad24 pre-Ad plasmid, composed entirely of Ad24 sequence and containing an E1 deletion and an E4 deletion substituted with Ad5 Orf6), an Ad24 ITR cassette was constructed containing sequences from the right (bp 31978 to 32264 and bp 34713 to 35164) and left (bp 4 to 450 and bp 3364 to 3799) end of the Ad24 genome separated by plasmid sequences containing a bacterial origin of replication and an ampicillin resistance gene. These four segments were generated by PCR and cloned sequentially into pNEB193, generating pNEBAd24-4. Next the Ad5 Orf6 open reading frame (Ad5 bp 31192 to bp 34078) was generated by PCR and cloned between Ad24 bp 32264 and 34713 generating

pNEBAd24E-Ad5Orf6 (the ITR cassette). PNEB193 is a commonly used commercially available cloning plasmid (New England Biolabs cat# N3051S) containing a bacterial origin of replication, ampicillin resistance gene and a multiple cloning site into which the PCR products were introduced. The ITR cassette contains a deletion of E1 sequences from Ad24 bp 451 to 3363 with a unique Swa I restriction site located in the deletion and an E4 deletion from Ad24 bp 32265 to 34712 into which Ad5 Orf6 was introduced in an E4 parallel orientation. In this 5 construct Ad5 Orf6 expression is driven by the Ad24 E4 promoter. The Ad24 sequences (bp 31978 to 32264 and bp 3464 to 3799) in the ITR cassette provide regions of homology with the purified Ad24 viral DNA in which bacterial recombination can occur following cotransformation into BJ 5183 bacteria (Figure 22). The ITR cassette was also designed to contain unique restriction enzyme sites (PmeI) located at the end of the viral ITR's so that digestion will release 10 the recombinant Ad24 genome from plasmid sequences. Potential clones will be screened by restriction analysis and one clone was selected as pMRKAd24ΔE1ΔE4Ad5Orf6. Pre-Adenovirus plasmid pMRKAd24ΔE1ΔE4Ad5Orf6 should contain Ad24 sequences from bp 4 to 450; bp 3364 to bp 32264 and bp 34713 to bp 35164 with Ad5Orf6 cloned between bp 32264 and bp 34713. The bp numbering in the above description refers to the wt sequence for both 15 Ad24 and Ad5.

EXAMPLE 16

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Insertion of HIV-1 gag and SEAP transgenes into pAd24ΔΕ1ΔΕ4Ad5Orf6 20

In order to introduce a gag or SEAP expression cassettes into the E1 region of pMRKAd24ΔE1ΔE4Ad5Orf6, bacterial recombination will be used. An HIV-1 gag expression cassette will consist of the following: 1) the immediate early gene promoter from the human cytomegalovirus, 2) the coding sequence of the human immunodeficiency virus type 1 (HIV-1) gag (strain CAM-1; 1526 bp) gene, and 3) the bovine growth hormone polyadenylation signal sequence, in the E1 deletion of an Ad24 shuttle plasmid, pNEBAd24-2 (a precursor to the Ad24 ITR cassette described above), generating pNEBAd24CMVgagBGHpA. PNEBAd24-2 contains Ad24 sequences from the left end of the genome (bp 4 to 450 and bp 3364 to 3799) that define the E1 deletion. The gag expression cassette will be obtained from a previously constructed plasmid and cloned into the E1 deletion between bp 450 and 3364 in the E1 parallel orientation. The shuttle vector containing the gag transgene will be digested to generate a DNA fragment consisting of the gag expression cassette flanked by Ad24 bp 4 to 450 and bp 3364 to 3799 and the fragment will be purified after electrophoresis on an agarose gel. Cotransformation of BJ 5183 bacteria with the shuttle vector fragment and pMRKAd24 Δ E1 Δ E4Ad5Orf6 which was linearized in the E1 region by digestion with SwaI, should result in the generation of Ad24 gag-

containing pre-Adenovirus plasmids pMRKAd24 Δ E1gag Δ E4Ad5Orf6 by homologous recombination. Potential clones will be screened by restriction analysis.

A similar strategy will be used to generate Ad24 pre-Ad plasmids containing a SEAP expression cassette. In this case, a SEAP expression cassette will consist of: 1) the immediate early gene promoter from the human cytomegalovirus, 2) the coding sequence of the human placental SEAP gene, and 3) the bovine growth hormone polyadenylation signal sequence cloned into the E1 deletion of an Ad24 shuttle plasmid, pNEBAd24-2, generating pNEBAd24CMVSEAPBGHpA. The transgene will then be recombined into pMRKAd24 Δ E1 Δ E4Ad5Orf6 as described above for the gag transgene.

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EXAMPLE 17

In Vivo Immunogenicity

A. Immunization

Rhesus macaques were between 3-10 kg in weight. In all cases, the total dose of each vaccine was suspended in 1 mL of buffer. The macaques were anesthetized (ketamine/xylazine) and the vaccines were delivered i.m. in 0.5-mL aliquots into both deltoid muscles using tuberculin syringes (Becton-Dickinson, Franklin Lakes, NJ). Peripheral blood mononuclear cells (PBMC) were prepared from blood samples collected at several time points during the immunization regimen. All animal care and treatment were in accordance with standards approved by the Institutional Animal Care and Use Committee according to the principles set forth in the *Guide for Care and Use of Laboratory Animals*, Institute of Laboratory Animal Resources, National Research Council.

25 B. T Cell Responses

Ad24 Vaccine Vector as a Heterologous Booster: Cohort of 4 rhesus macaques was immunized initially with 3 doses (wk 0, 4, 26) of either 10⁷ or 10⁹ vp of MRKAd5-gag (see, PCT/US01/28861, published March 21, 2002) or MRKAd6-gag. At wk 56, the animals received a booster vaccine of 10¹¹ vp Ad24ΔE1gagΔOrf6Ad5Orf6. A separate cohort of naïve animals received a single dose of the booster vaccine. The results of the IFN-γ ELISPOT analyses of PBMC collected during the course of the studies are shown in Figure 23. It is apparent that the Ad24 HIV vectors can be utilized to amplify the existing pools of HIV-specific T cells. The increases in the levels of gag-specific T cells from the pre-boost levels to those measured at 4 wks post boost were consistently larger than the levels induced by the same booster vaccine in naïve animals. PBMCs from the vaccinees of the heterologous MRKAd5/MRKAd6-Ad24 boost

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regimen were analyzed for intracellular IFN-γ staining after the priming immunizations (wk 60). The assay results provided information on the relative amounts of CD4⁺ and CD8⁺ gag-specific T cells in the peripheral blood (Figure 24). The results indicated that heterologous prime-boost immunization approach was able to elicit in rhesus macaques both HIV-specific CD4+ and CD8+ T cells.

Ad24 Vaccine Vector as a Heterologous Primer: In a separate study, a cohort of 3 rhesus macaques was immunized initially with 2 doses (wk 0, 4) of 10¹¹ vp Ad24ΔE1gagΔOrf6Ad5Orf6 and boosted at wk 24 with 10⁷ vp of MRKAd5-gag. The low dose of MRKAd5-gag is selected to mimic the effect of pre-existing neutralizing immunity to the vector in a subject. A separate cohort of naïve animals was given a single dose of 10⁷ vp MRKAd5-gag. The results of the IFN- γ ELISPOT analyses of PBMC collected during the course of the studies are shown in Figure 25.

The Ad24-based vaccine was able to prime effectively for HIV-specific T cell responses in macaques. Boosting with a low dose MRKAd5-gag resulted in a significant increase in the levels of gag-specific T cells. The increases in 2 out of 3 animals exceed the levels typically observed after treatment of naïve animals with the same low dose of MRKAd5gag.

EXAMPLE 18

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Construction of pAd34AE1AE4Ad5Orf6 20

To generate an E1- Ad34 based vector that can propagate in existing group C/Ad5 E1 complementing cell lines (293, PER.C6), Ad5 Orf6 was inserted in place of the native E4 region. Since at the time, the complete sequence of Ad34 (see Figures 28A-1 to 28A-9; subject of copending application serial no. 60/458,825, filed March 28, 2003) was unknown, advantage was taken of the sequence homology between Ad34 and Ad35 in order to construct the Ad34 pre-Adenovirus plasmid. Cotransformation of BJ 5183 bacteria with purified wild-type Ad34 viral DNA and the appropriately constructed Ad35 ITR cassette resulted in the circularization of the viral genome by homologous recombination. The construction of the pre-Ad plasmid based on Ad34, is outlined below:

To construct pAd34ΔE1ΔE4Ad5Orf6 (An Ad34 pre-Ad plasmid containing an E1 deletion and an E4 deletion substituted with Ad5 Orf6), we utilized an Ad35 ITR cassette. We anticipated that sequence homology between Ad34 and Ad35 would allow homologous recombination to occur. The Ad35 ITR cassette was constructed containing sequences from the right (bp 31599 to 31913 and bp 34419 to 34793) and left (bp 4 to 456 and bp 3403 to 3886) end of the Ad35 genome (see Figures 2A-1 to 2A-10) separated by plasmid sequences containing a

bacterial origin of replication and an ampicillin resistance gene. The four segments were generated by PCR and cloned sequentially into pNEB193, generating pNEBAd35-4. Next the Ad5 Orf6 open reading frame was generated by PCR and cloned between Ad35 bp 31913 and 34419 generating pNEBAd35-4Ad5Orf6 (the ITR cassette). PNEB193 is a commonly used commercially available cloning plasmid (New England Biolabs cat# N3051S) containing a bacterial origin of replication, ampicillin resistance gene and a multiple cloning site into which the PCR products were introduced. The ITR cassette contains a deletion of E1 sequences from Ad35 bp 457 to 3402 with a unique Swa I restriction site located in the deletion and an E4 deletion from Ad35 bp 31914 to 34418 into which Ad5 Orf6 was introduced in an E4 parallel orientation. In this construct Ad5Orf6 expression is driven by the Ad35 E4 promoter. The Ad35 sequences (bp 31599 to 31913 and bp 3403 to 3886) in the ITR cassette provided regions of homology with the purified Ad34 viral DNA in which bacterial recombination could occur following cotransformation into BJ 5183 bacteria (Figure 26). The ITR cassette was also designed to contain unique restriction enzyme sites (PmeI) located at the end of the viral ITR's so that digestion would release the recombinant Ad34 genome from the plasmid sequences. Potential clones were screened by restriction analysis and one clone was selected as pAd 34Δ E 1Δ E4Ad5Orf6.

EXAMPLE 19

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20 Rescue of pAd34ΔΕ1ΔΕ4Ad5Orf6 into Virus

In order to determine if pre-adenovirus plasmid pAd34AE1AE4Ad5Orf6, could be rescued into virus and propagated in a group C E1 complementing cell line, the plasmid was digested with *Pme* I and transfected into T-25 flasks of PER.C6 cells using the calcium phosphate co-precipitation technique (Cell Phect Transfection Kit, Amersham Pharmacia Biotech Inc). *Pme*I digestion releases the viral genome from plasmid sequences allowing viral replication to occur after cell entry. Viral cytopathic effect (CPE), indicating that virus replication and amplification was occurring was observed following transfection. When CPE was complete, approximately 7-10 days post transfection, the infected cells and media were harvested, freeze/thawed three times and the cell debris pelleted by centrifugation.

Approximately 1 ml of the cell lysate was used to infect a T-225 flask of PER.C6 cells at 80-90% confluence. Once CPE was reached, infected cells and media were harvested, freeze/thawed three times and the cell debris pelleted by centrifugation. Clarified cell lysates were then used to infect 2-layer NUNC cell factories of PER.C6 cells. Following complete CPE, the virus was purified by ultracentrifugation on CsCl density gradients. In order to verify the genetic structure of the rescued viruses, viral DNA was extracted using pronase treatment

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followed by phenol chloroform extraction and ethanol precipitation. Viral DNA was then digested with HindIII and treated with Klenow fragment to end-label the restriction fragments with P33-dATP. The end-labeled restriction fragments were then size-fractionated by gel electrophoresis and visualized by autoradiography. The digestion products were compared with the digestion products of the corresponding pre-Adenovirus plasmid (that had been digested with Pme1/HindIII prior to labeling) from which they were derived. The expected sizes were observed, indicating that the viruses had been successfully rescued.

EXAMPLE 20

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Insertion of an Expression Cassette into pAd34ΔE1ΔE4Ad5Orf6

In order to introduce a gag or SEAP expression cassette (see Figures 6 and 7, respectively) into the E1 region of pAd34ΔE1ΔE4Ad5Orf6, bacterial recombination was again used. A gag expression cassette consisting of the following: 1) the immediate early gene promoter from human cytomegalovirus, 2) the coding sequence of the human immunodeficiency virus type 1 (HIV-1) gag (strain CAM-1; 1526 bp) gene, and 3) the bovine growth hormone polyadenylation signal sequence, was cloned into the E1 deletion in Ad35 shuttle plasmid, pNEBAd35-2 (a precursor to the Ad35 ITR cassettes described above), generating pNEBAd35CMVgagBGHpA. pNEBAd35-2 contains Ad35 sequences from the left end of the genome (bp 4 to 456 and bp 3403 to 3886) with a unique SwaI site between bp 456 and 3403 at the position of the deletion. The gag expression cassette was obtained from a previously constructed shuttle plasmid by EcoRI digestion. Following the digestion the desired fragment was gel purified, treated with Klenow to obtain blunt ends and cloned into the SwaI site in pNEBAd35-2. This cloning step resulted in the gag expression cassette being inserted into the E1 deletion between bp 456 and 3403 in the E1 parallel orientation. The shuttle vector containing the gag transgene was digested to generate a DNA fragment consisting of the gag expression cassette flanked by Ad35 bp 4 to 456 and bp 3403 to 3886 and the fragment was purified after electrophoresis on an agarose gel. Cotransformation of BJ 5183 bacteria with the shuttle vector fragment and pAd34 Δ E1 Δ E4Ad5Orf6, linearized in the E1 region by digestion with Swa I, resulted in the generation of the Ad34 gag-containing pre-Adenovirus plasmid pAd34ΔE1gagΔE4Ad5Orf6 by homologous recombination. Potential clones were screened by 30 restriction analysis.

A similar strategy was used to generate Ad34 pre-Ad plasmids containing a SEAP expression cassette. In this case a SEAP expression cassette consisting of: 1) the immediate early gene promoter from human cytomegalovirus, 2) the coding sequence of the human placental SEAP gene, and 3) the bovine growth hormone polyadenylation signal sequence was

cloned into the E1 deletion in Ad35 shuttle plasmid, pNEBAd35-2, generating pNEBAd35CMVSEAPBGHpA. The SEAP expression cassette was obtained from a previously constructed shuttle plasmid by *Eco*RI digestion. Following the digestion the desired fragment was gel purified, treated with Klenow to obtain blunt ends and cloned into the *Swa*I site in pNEBAd35-2. The transgene was then recombined into the pAd34ΔE1ΔE4Ad5Orf6, generating pAd34ΔE1SEAPΔE4Ad5Orf6 as described above for the gag transgene.

All pre-Ad plasmids were rescued into virus and expanded to prepare CsCl purified stocks as described above.

EXAMPLE 21

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Construction of pMRKAd34ΔE1ΔE4Ad5Orf6

To construct an Ad34 pre-Ad plasmid that was composed entirely of Ad34 sequences, an Ad34 ITR cassette was generated. The Ad34 ITR cassette was constructed containing sequences from the right (bp 31584 to 31895 and bp 34409 to 34772) and left (bp 4 to 456 and bp 3402 to 3885) end of the Ad34 genome (see Figures 28A-1 to 28A-9) separated by 15 plasmid sequences containing a bacterial origin of replication and an ampicillin resistance gene. These four segments were generated by PCR and cloned sequentially into pNEB193, generating pNEBAd34-4. Next the Ad5 Orf6 open reading frame was generated by PCR and cloned between Ad34 bp 31895 and 34409 generating pNEBAd34-4Ad5Orf6 (the ITR cassette). PNEB193 is a commonly used commercially available cloning plasmid (New England Biolabs 20 cat# N3051S) containing a bacterial origin of replication, ampicillin resistance gene and a multiple cloning site into which the PCR products were introduced. The ITR cassette contains a deletion of E1 sequences from Ad34 bp 457 to 3401 with a unique Swa I restriction site located in the deletion and an E4 deletion from Ad34 bp 31896 to 34408 into which Ad5 Orf6 was introduced in an E4 parallel orientation. In this construct Ad5Orf6 expression is driven by the 25 Ad34 E4 promoter. The Ad34 sequences (bp 31584 to 31895 and bp 3402 to 3885) in the ITR cassette provided regions of homology with the purified Ad34 viral DNA in which bacterial recombination could occur following cotransformation into BJ 5183 bacteria (Figure 27). The ITR cassette was also designed to contain unique restriction enzyme sites (PmeI) located at the end of the viral ITR's so that digestion would release the recombinant Ad34 genome from the 30 plasmid sequences. Potential clones were screened by restriction analysis and one clone was selected as pMRKAd34ΔE1ΔE4Ad5Orf6.

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EXAMPLE 22 In Vivo Studies

A. Immunization

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Cohorts of 3 rhesus macaques were given single intramuscular injections of one of the two vectors: (1) 10^11 vp MRKAd5-SEAP (in MRKAd vector backbone disclosed in PCT/US01/28861, published March 21, 2002); and (2) 10¹¹ vp Ad34ΔE1SEAPΔE4Ad5Orf6. Rhesus macaques were between 3-10 kg in weight. In all cases, the total dose of each vaccine was suspended in 1 mL of buffer. The macaques were anesthetized (ketamine/xylazine) and the vaccines were delivered i.m. in 0.5-mL aliquots into both deltoid muscles using tuberculin syringes (Becton-Dickinson, Franklin Lakes, NJ). Peripheral blood mononuclear cells (PBMC) were prepared from blood samples collected at several time points during the immunization regimen. All animal care and treatment were in accordance with standards approved by the Institutional Animal Care and Use Committee according to the principles set forth in the Guide for Care and Use of Laboratory Animals, Institute of Laboratory Animal Resources, National Research Council.

B. SEAP Assay

Serum samples were analyzed for circulating human secreted alkaline phosphatase (SEAP) levels using TROPIX phospha-light chemiluminescent kit (Applied Biosystems Inc). Duplicate 5 μ L aliquots of each serum were mixed with 45 μ L of kit-supplied dilution buffer in a 96-well white DYNEX plate. Serially diluted solutions of a human placental alkaline phosphatase (Catalog no. M5905, Sigma, St. Louis, MO) in 10% naïve monkey serum served to provide the standard curve. Endogenous SEAP activity in the samples was inactivated by heating the well for 30 minutes at 65 °C. Enzymatic SEAP activities in the samples were determined following the procedures described in the kit. Chemiluminescence readings (in relative light units) were recorded using DYNEX luminometer. RLU readings were converted to ng/mL SEAP using a log-log regression analyses.

C. ELISPOT Assay 30

The IFN-γ ELISPOT assays for rhesus macaques were conducted following a previously described protocol (Allen et al., 2001 J. Virol. 75(2):738-749), with some modifications. For antigen-specific stimulation, a peptide pool was prepared from 20-aa peptides that encompass the entire HIV-1 gag sequence with 10-aa overlaps (Synpep Corp., Dublin, CA). To each well, 50 μ L of 2-4 x 10^5 peripheral blood mononuclear cells (PBMCs) were added; the cells were counted using Beckman Coulter Z2 particle analyzer with a lower

size cut-off set at 80 femtoliters ("fL"). Either 50 μ L of media or the gag peptide pool at 8 μ g/mL concentration per peptide was added to the PBMC. The samples were incubated at 37°C, 5% CO₂ for 20-24 hrs. Spots were developed accordingly and the plates were processed using custom-built imager and automatic counting subroutine based on the ImagePro platform (Silver Spring, MD); the counts were normalized to 10^6 cell input.

D. Intracellular Cytokine Staining (ICS)

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To 1 ml of 2 x 106 PBMC/mL in complete RPMI media (in 17x100mm round bottom polypropylene tubes (Sarstedt, Newton, NC)), anti-hCD28 (clone L293, Becton-Dickinson) and anti-hCD49d (clone L25, Becton-Dickinson) monoclonal antibodies were added to a final concentration of 1 μ g/mL. For gag-specific stimulation, 10 μ L of the peptide pool (at 0.4 mg/mL per peptide) were added. The tubes were incubated at 37 °C for 1 hr., after which 20 μL of 5 mg/mL of brefeldin A (Sigma) were added. The cells were incubated for 16 hr at 37 °C, 5% CO₂, 90% humidity. 4 mL cold PBS/2%FBS were added to each tube and the cells were pelleted for 10 min at 1200 rpm. The cells were re-suspended in PBS/2%FBS and stained (30 min, 4 °C) for surface markers using several fluorescent-tagged mAbs: 20 μL per tube antihCD3-APC, clone FN-18 (Biosource); 20 μL anti-hCD8-PerCP, clone SK1 (Becton Dickinson); and 20 µL anti-hCD4-PE, clone SK3 (Becton Dickinson). Sample handling from this stage was conducted in the dark. The cells were washed and incubated in 750 μ L 1xFACS Perm buffer (Becton Dickinson) for 10 min at room temperature. The cells were pelleted and re-suspended in PBS/2%FBS and 0.1 μg of FITC-anti-hIFN-γ, clone MD-1 (Biosource) was added. After 30 min incubation, the cells were washed and re-suspended in PBS. Samples were analyzed using all four color channels of the Becton Dickinson FACSCalibur instrument. To analyze the data, the low side- and forward-scatter lymphocyte population was initially gated; a common fluorescence cut-off for cytokine-positive events was used for both CD4⁺ and CD8⁺ populations, and for both mock and gag-peptide reaction tubes of a sample.

E. Results

Expression: Serum samples prior to and after the injection were analyzed for circulating SEAP activities and the results are shown in Figure 29. Results indicate that the peak levels of SEAP protein produced by the alternative adenovirus serotype were lower than but were within 3-fold of that of MRKAd5 at the same high dose level of 10^11 vp (Figure 29). The levels of SEAP in the serum dropped dramatically after day 10 and were close to background as early as day 15. These observations strongly indicate that the Ad34-based vector is efficient in expressing a transgene following intramuscular administration in a primate.

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Immunogenicity: Vaccine-induced T cell responses against HIV-1 gag were quantified using IFN-gamma ELISPOT assay against a pool of 20-aa peptides that encompassed the entire protein sequence. The results are shown in Figure 30; they are expressed as the number of spot-forming cells (SFC) per million peripheral blood mononuclear cells (PBMCs) that responded to the peptide pool or the mock (no peptide) control.

Immunization with gag-expressing Ad34 vector induced detectable levels of circulating gag-specific T cells immediately after a single dose of the vector. The responses improved following a second dose given at wk 4. Overall, the responses to the Ad34-based vector were slightly lower than those induced by the same dose of MRKAd5-gag. The results strongly indicate the Ad34-based vector can prime effectively for HIV-specific T cell responses.

IFN-γ ICS analyses of the PBMC from the Ad34-immunized animals revealed that the vector can induce detectable levels of both CD4⁺ and CD8⁺ HIV-specific T cells (Figure 31).

EXAMPLE 23 15

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Heterologous Immunization

Cohorts of 3 monkeys were immunized (at wks 0, 4) with 10^11 vp Ad34ΔE1gagΔE4Ad5Orf6 followed by a booster at week 24 with 10^10 vp Ad35ΔE1gagΔE4Ad5Orf6. Vaccine-induced T cell responses against HIV-1 gag were quantified using IFN-gamma ELISPOT assay against a pool of 20-aa peptides that encompassed the entire protein sequence. The results are shown in Figure 32; they are expressed as the number of spot-forming cells (SFC) per million peripheral blood mononuclear cells (PBMCs) that responded to the peptide pool or the mock (no peptide) control.

Immunization with gag-expressing Ad34 vector induced detectable levels of circulating gag-specific T cells that decreased to between 94-139 SFC/10^6 PBMC at the time of the boost. Heterologous immunization with an Ad35-based HIV vector resulted in as much as a 3-fold increase in T cell responses.

IFN-γ ICS analyses of the PBMCs from the Ad34 primed/Ad35 boosted animals at week 28 revealed that the vector can induce detectable levels of both CD4+ and CD8+ HIVspecific T cells (Figure 33).

WHAT IS CLAIMED IS:

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1. A means for propagating replication-defective adenovirus in an adenoviral E1-complementing cell line expressing E1 gene product(s) which are non-native to the adenovirus, which comprises:

- (a) inserting all or a portion of a heterologous adenoviral E4 region comprising nucleic acid sequence encoding open reading frame 6 (ORF6) into a replication-defective adenovirus; wherein the E4 region or portion thereof inserted into the adenovirus is native to a virus of the same adenovirus serotype as the E1 gene product(s) expressed by the complementing cell line;
- (b) introducing the replication-defective adenovirus into the adenoviral E1-complementing cell line;
- (c) allowing the replication-defective adenovirus to propagate in the adenoviral E1-complementing cell line; and
 - (d) rescuing the propagated adenovirus.
- 2. A means in accordance with claim 1 wherein the heterologous adenoviral E4 region or portion thereof comprises the complete adenoviral E4-encoding region.
- 3. A means in accordance with claim 2 wherein the heterologous adenoviral E4 region or portion thereof comprises the complete adenoviral E4-encoding region and native E4 promoter.
- 4. A means in accordance with claim 1 wherein the heterologous adenoviral E4 region or portion thereof is inserted into the replication-defective virus in place of nucleic acid sequence encoding open reading frame 6 (ORF6).

5. A means in accordance with claim 1 wherein the heterologous adenoviral E4 region or portion thereof is inserted into the replication-defective virus in place of nucleic acid sequence encoding the complete adenoviral E4-encoding region.

- 6. A means in accordance with claim 1 wherein the heterologous adenoviral
 5 E4 region or portion thereof is derived from a subgroup C adenovirus.
 - 7. A means in accordance with claim 1 wherein the subgroup C adenovirus is adenovirus of serotype 5.
 - 8. A means in accordance with claim 7 wherein the replication-defective adenovirus is an adenovirus of subgroup B.
 - A means in accordance with claim 7 wherein the replication-defective adenovirus is an adenovirus of serotype 35.

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- 10. A means in accordance with claim 1 wherein the heterologous adenoviralE4 region or portion thereof is operatively linked to a heterologous promoter.
- 11. A means in accordance with claim 1 wherein the adenoviral E1-complementing cell line is a PER.C6® cell line.
 - 12. A replication-defective adenovirus comprising all or a portion of a heterologous E4 region comprising a heterologous adenoviral open reading frame 6 (ORF6).
 - 13. A replication-defective adenovirus in accordance with claim 12 wherein the adenovirus comprises a heterologous gene of interest.
- 14. A replication-defective adenovirus in accordance with claim 13 wherein the heterologous gene of interest is a gene encoding an HIV-1 antigen.
 - 15. A replication-defective adenovirus in accordance with claim 14 wherein the HIV-1 antigen is selected from the group consisting of HIV-1 gag, pol, nef and env.

16. A replication-defective adenovirus comprising all or a portion of a heterologous E4 region comprising a heterologous adenoviral open reading frame 6 (ORF6) and a gene encoding HIV-1 gag.

- 17. A replication-defective adenovirus comprising all or a portion of a

 beterologous E4 region comprising a heterologous adenoviral open reading frame 6 (ORF6) in

 place of a native E4 region or portion thereof comprising ORF6.
 - 18. A replication-defective adenovirus comprising all or a portion of a heterologous E4 region comprising a complete heterologous E4 region in place of a complete native E4 region.
 - 19. A replication-defective adenovirus comprising a heterologous E4 region or portion thereof comprising a complete heterologous E4 region including E4 promoter in place of a complete native E4 region.

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- 20. Adenovirus propagated in accordance with the means of claim 1.
- 21. A means in accordance with claim 1 wherein the replication-defective adenovirus comprises a heterologous gene of interest.
 - 22. A means in accordance with claim 21 wherein the heterologous gene of interest is a gene encoding an HIV-1 antigen.
 - 23. A means in accordance with claim 22 wherein the HIV-1 antigen is selected from the group consisting of: HIV-1 gag, pol, nef and env.
 - 24. A replication-defective adenovirus of serotype 35 comprising all or a portion of an adenovirus serotype 5 E4 region comprising open reading frame 6 (ORF6) and a heterologous gene of interest.
 - 25. A replication-defective adenovirus in accordance with claim 24 wherein the heterologous gene of interest is a gene encoding an HIV-1 antigen.

26. A replication-defective adenovirus in accordance with claim 25 wherein the HIV-1 antigen is selected from the group consisting of: HIV-1 gag, pol, nef and env.

27. A replication-defective adenovirus of serotype 35 comprising all or a portion of an adenovirus serotype 5 E4 region comprising open reading frame 6 (ORF6) and a gene encoding HIV-1 gag.

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- 28. A recombinant adenoviral vector of serotype 24 which comprises an E4 gene or a segment of an E4 gene comprising open reading frame 6 ("ORF6") of an alternative serotype.
- 29. A population of cells comprising the recombinant adenoviral vector of10 claim 28.
 - 30. A method for producing recombinant, replication-defective adenovirus particles comprising:
 - (a) introducing a recombinant adenoviral vector of claim 28 into a population of cells expressing adenovirus E1; and
 - (b) harvesting the resultant recombinant, replication-defective adenovirus.
 - 31. Purified recombinant, replication-defective adenovirus particles harvested in accordance with the method of claim 30.
 - 32. A composition comprising purified recombinant adenovirus particles in accordance with claim 31.
- 20 33. A composition in accordance with claim 32 which comprises a physiologically acceptable carrier.
 - 34. A recombinant adenoviral vector in accordance with claim 28 which is at least partially deleted in E1 and devoid of E1 activity and comprises a heterologous nucleic acid.

35. A composition comprising purified recombinant adenoviral particles in accordance with claim 31 which are at least partially deleted in E1 and devoid of E1 activity and comprise a heterologous nucleic acid.

36. A method for effecting the delivery and expression of heterologous nucleic acid comprising administering the composition of claim 35 prior or subsequent to administration of the heterologous nucleic acid with the same or different vector.

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- 37. A method in accordance with claim 36 wherein the composition is preceded or followed by administration of heterologous nucleic acid with an adenovirus of a different serotype.
- 38. A composition in accordance with claim 35 wherein the heterologous nucleic acid encodes an HIV antigen.
- 39. A method for generating a cellular-mediated immune response against HIV in an individual comprising administering to the individual a composition of claim 38.
- 40. A composition in accordance with claim 39 wherein the HIV antigen is

 HIV-1 gag or immunologically relevant modification thereof.
 - 41. A composition in accordance with claim 39 wherein the HIV antigen is HIV-1 nef or immunologically relevant modification thereof.
 - 42. A composition in accordance with claim 39 wherein the HIV antigen is HIV-1 pol or immunologically relevant modification thereof.
- 43. A recombinant adenoviral vector of serotype 24 which is at least partially deleted in E1 and devoid of E1 activity; wherein said vector comprises an E4 gene or a segment of an E4 gene from adenovirus serotype 5 comprising open reading frame 6 ("ORF6"), and a heterologous nucleic acid.

44. A population of cells comprising the recombinant adenoviral vector of claim 43.

45. A method for producing recombinant, replication-defective adenovirus particles comprising:

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- (a) introducing a recombinant adenoviral vector of claim 43 into a population of cells expressing adenovirus serotype 5 E1; and
 - (b) harvesting the resultant recombinant, replication-defective adenovirus.
- 46. Purified recombinant, replication-defective adenovirus particles harvested in accordance with the method of claim 45.
- 47. A composition comprising purified recombinant adenovirus particles in accordance with claim 46.
- 48. A composition in accordance with claim 47 which comprises a physiologically acceptable carrier.
- 49. A method for effecting the delivery and expression of the heterologous nucleic acid comprising administering the composition of claim 48 prior or subsequent to administration of the heterologous nucleic acid with the same or different vector.
- 50. A method in accordance with claim 49 above wherein the composition is preceded or followed by administration of the heterologous nucleic acid with an adenovirus of a different serotype.
- 51. A composition in accordance with claim 48 wherein the heterologous nucleic acid encodes an HIV antigen.
- 52. A method for generating a cellular-mediated immune response against HIV in an individual comprising administering to the individual a composition of claim 51.

53. A composition in accordance with claim 51 wherein the HIV antigen is HIV-1 gag or immunologically relevant modification thereof.

- 54. A composition in accordance with claim 51 wherein the HIV antigen is HIV-1 nef or immunologically relevant modification thereof.
- 55. A composition in accordance with claim 51 wherein the HIV antigen is HIV-1 pol or immunologically relevant modification thereof.

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- A recombinant adenoviral vector of serotype 34 which comprises an E4 gene or a segment of an E4 gene comprising open reading frame 6 ("ORF6") of an alternative serotype.
- 57. A population of cells comprising the recombinant adenoviral vector of claim 56.
- 58. A method for producing recombinant, replication-defective adenovirus particles comprising:
- (a) introducing a recombinant adenoviral vector of claim 56 into a population of cells expressing adenovirus E1; and
 - (b) harvesting the resultant recombinant, replication-defective adenovirus.
 - 59. Purified recombinant, replication-defective adenovirus particles harvested in accordance with the method of claim 58.
- 60. A composition comprising purified recombinant adenovirus particles in accordance with claim 59.
 - 61. A composition in accordance with claim 60 which comprises a physiologically acceptable carrier.
 - 62. A recombinant adenoviral vector in accordance with claim 56 which is at least partially deleted in E1 and devoid of E1 activity and comprises a heterologous nucleic acid.

63. A composition comprising purified recombinant adenoviral particles in accordance with claim 59 which are at least partially deleted in E1 and devoid of E1 activity and comprise a heterologous nucleic acid.

64. A method for effecting the delivery and expression of heterologous nucleic acid comprising administering the composition of claim 63 prior or subsequent to administration of the heterologous nucleic acid with the same or different vector.

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- 65. A method in accordance with claim 64 wherein the composition is preceded or followed by administration of heterologous nucleic acid with an adenovirus of a different serotype.
- 10 66. A composition in accordance with claim 63 wherein the heterologous nucleic acid encodes an HIV antigen.
 - 67. A method for generating a cellular-mediated immune response against HIV in an individual comprising administering to the individual a composition of claim 66.
 - 68. A composition in accordance with claim 67 wherein the HIV antigen is HIV-1 gag or immunologically relevant modification thereof.
 - 69. A composition in accordance with claim 67 wherein the HIV antigen is HIV-1 nef or immunologically relevant modification thereof.
 - 70. A composition in accordance with claim 67 wherein the HIV antigen is HIV-1 pol or immunologically relevant modification thereof.
 - 20 71. A recombinant adenoviral vector of serotype 34 which is at least partially deleted in E1 and devoid of E1 activity; wherein said vector comprises an E4 gene or a segment of an E4 gene from adenovirus serotype 5 comprising open reading frame 6 ("ORF6"), and a heterologous nucleic acid.

72. A population of cells comprising the recombinant adenoviral vector of claim 71.

- 73. A method for producing recombinant, replication-defective adenovirus particles comprising:
- (a) introducing a recombinant adenoviral vector of claim 71 into a population of cells expressing adenovirus serotype 5 E1; and

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- (b) harvesting the resultant recombinant, replication-defective adenovirus.
- 74. Purified recombinant, replication-defective adenovirus particles harvested in accordance with the method of claim 73.
- 75. A composition comprising purified recombinant adenovirus particles in accordance with claim 74.
- 76. A composition in accordance with claim 75 which comprises a physiologically acceptable carrier.
- 77. A method for effecting the delivery and expression of the heterologous nucleic acid comprising administering the composition of claim 76 prior or subsequent to administration of the heterologous nucleic acid with the same or different vector.
- 78. A method in accordance with claim 77 above wherein the composition is preceded or followed by administration of the heterologous nucleic acid with an adenovirus of a different serotype.
- 79. A composition in accordance with claim 76 wherein the heterologous nucleic acid encodes an HIV antigen.
- 80. A method for generating a cellular-mediated immune response against HIV in an individual comprising administering to the individual a composition of claim 79.

81. A composition in accordance with claim 79 wherein the HIV antigen is HIV-1 gag or immunologically relevant modification thereof.

- 82. A composition in accordance with claim 79 wherein the HIV antigen is HIV-1 nef or immunologically relevant modification thereof.
- 83. A composition in accordance with claim 79 wherein the HIV antigen is HIV-1 pol or immunologically relevant modification thereof.

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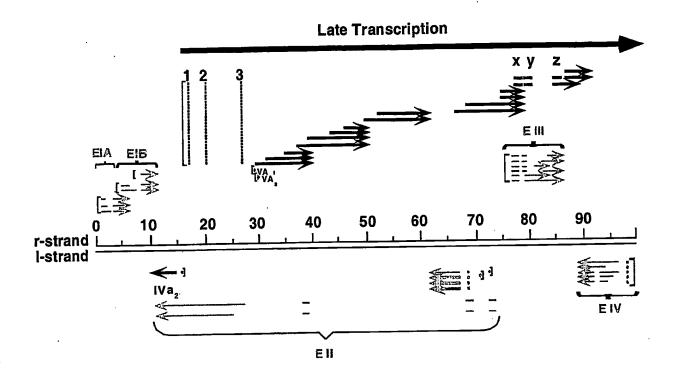


FIG. 1

_		t-t-cott	atagatggaa	tagtgccaat	atgtaaatga g gttaaaaggg 9	gtgatttta
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61	aaaagtgtgg	geegrgrygr	taaaaataaa	attttttgc	aagttgtcgc g ttcccacggt a	ggaaatgtt
121	cgtgggaaaa	tgacgillia	tataacaaa	ctacttagtt	ttcccacggt a attttcgcgc g	tttaacagg
181	acgcataaaa	aggettett	CCCCCGGGGG	anattacta	attttcgcgc 9	_{raaaactgaa}
241	aaatgaggta	gttttgaccy	tactataata	tttatggcag	ggtggagtat t	tgttcaggg
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37	21 ctatggaag	ge accoraged				

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	780T	granager	ggatttcctg	ccaccagttg	gaggattggc	tgttgatgtg a acagacggcc g	atggaagtag
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28981	tgacttctgo	tegeteacae	beetersons	tatatggcoo	tagatagttt	acaaaaccat
29041	. ctcaaggtgg	tcatgtcttt	tggtggagaa	catatgatat	aaccattato	aacqtqacaq
29101	. gtgaccaacc	tggtagattt	ttetgeaacg	gcayayaccc	tagtttagat	tataacatta
		, <u>ammettetat</u>	rarodaaccu	actatadag		
29221	ttgtactgcc	atctaccact	ccagcacccc	geacaactac	coccage	atastast
			00000000000	reactions	. addacucuc	gcgaacaa
		- nantanaaa	atttccactt	Caacaattat	Callatige	gcagogacaa
			********	CCTACLACUL	: CLUCLUCUAL	agamagaca
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		- tataasattt	- ++acttttat	Lactiucati	, Lucutatyta	gcacageeeg
			· aacttctada	CEGUALCEL	. ulucqaalig	CCCGCCCGCG
		+	accaaaatat	cocoocacu	. Cilauacica	LLLAAAACCA
			. +++++actfa	' CALLUCLLU	: Clacucty	, ccaacccag
		~ +-ataaaaa	. ~==~=~~	naaaaluca	i allucaacaa	Cogoggoode
2988	L ctgcctata	g tactccacca g ctatcgagaa	. yaatattii	tececess	tttaataato	attoctogaa
2994	l ttcttgctt	g ctatcgagaa	adalCayddo	· cattttta	ataccccta	tttgattttg
3000	l taattaata	t aatctgttgc	accataatt	. cattteryal	. acacccccc	attoccoca
3006	1 gctggaatg	c tcccaatgca	catgatcato	: cacaayaccc	. ayayyaacac	ccccacter
0010		a acatocaats	acactaata:	T ALLACUADA	ı Luaattata	t ccccaceac
3018	1 tccctgcta	t tagttactto	aacctaacc	g gcggagacga	a Cigadacact	. Lactacette

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30241 aatteegeeg aggatetget egatatggae g	roccocatet cagaacaacg acttgcccaa
30241 aattccgccg aggatctgct cgatatggac g 30301 ctacgcatcc gccagcagca ggaacgcgtg g	rccaaagage teagagatgt catecaaatt
shall atacacated accadeages agains .	acastate ctacdadate
anaki daccaardca dddddyycuc mar-a-a-	ascracasas attitutive
20021 accordacto accarregació	warstartsa dodiliquali
20101 startagaa LCaacccca as	
SAENI ASAFRATTALI UCUBLICIONI USTISIS	
aneni aggracette taccaatgaa toa	ctcacttccc tcttcccaac
20661 Carcaataad Glocolgous dans	the tage tactttaggg dodgetulde
anable talle talactors	Latet cffccanat daccadyaga
20701 attitagete etelectice	
angli atcongctca glyacitett caasis	
30001 caccettea tadacecayy goodan	and account that acadelada
20061 offictactt taddatytte date	tarana catacolor
21021 Atamagada dacilacase saccase	The coattorian togattayaa
21001 acadcaccca Llactadada tado	Tabanat tagaatttaa caacuulgac
211/1 actcassaca atdadctacy cyclamors	The standard to accept accept
21201 affforatad dygalageac care	ctttagtaft agtadaga
21261 casafford addacactur out	The section of the se
21221 MARKOCEEG ELAGEGGECA COSTO	- LL-L-LL Ltgactcttc todaaalcua
21201 Fracacada agadageada carre	Thereare astottotac accodicage
21//1 Francedadd ddicagacco dallan	
21EN1 GARACEGEAG CCAGCAGCAG CG	actacatoac tagttalyac
21561 actactaggg atagggaaaa coacaa	TT googtatoat tecetodau
21621 agaagtetat titeettigad tutti	streathfree agaaagcade
21601 attacctate coaldcade coados	thacadaada cdacaactaa
217/1 atacctaccc todecacate cooling	antagttatt ttgcctccac
21001 aataaagttt adguguutuu duudaan	and another and attiggated
21061 officially datagance decides	ti agtitcadad cdadccadic
21021 Cattagagat addCattgtt cought	topaggggtt tcacagtcca
21001 Formotcage datagatada datasas	the standard accordingly
22011 actortorod atolyactic 9949	Liberta aparcharaa dcadccdcty
22101 atcataatcc gadaacggca cogges	
20161 FORGOTOGO ECCULUCUA COMPANIA	The same accordance and accordance
22221 Maffitaata decellades seems	atattatta ataaaccata
22201 Frenchcada CottlyCage aggettern	i
22211 affagaaged Cleedgeeda access	The same agartarrea catalalyal
22/01 ccapacttta ataladatta dargares	the transfer and cancett antiatical
20161 CHOFFEEDEC ALGUGECTE COMMENT	
22521 GCAACCCAAL ALAACCCCCC 9944	tectetedae edidadedae
22501 santgaaccc tgctgattac daugattac	
226/1 Francatoa dadatateta 555555	at a company of CEEGCaudac
22701 sattitaac tooloayyac coas	acactatora tautcatagu
22761 agtasagetg geagadaag gaaga	' actogaattt cattiticit
22021 atracastot ducadeageg 99095	Total and address and a transfer and
2001 acaacutuut dattugggtto taara	The second of th
20011 GCGCAACCEE QUUALGALGG GGCGG	. T accettacco tottcccqtqt
22001 CAUCCEDOC AVARCACACA COSTITUTO	
sanci getanticaa ulacayoodo dooroo	Lasta daggetagea tatgeaadu
22121 transactor allegeaters accept	
22101 CCAACCAAGC AACGCAACCA STEEL	tosastrora dallucucucay
22211 GARCCATOET Adilities Courses	cotasatosa sadaaatucu
22201 atggcatete tegeceede egegees	a accordant ccaadacad
22261 atttcaagg tgctcaacgg tggccca	
22/21 samafacca dadyadygay Culture	cototoantt cttotootaa
22/01 Cattccada Ladicillay Colors	tacaccacca trettadaca
225/1 atroaatoca cacattacaa acussi	stancoatt gagaatggca
33601 cacceteata algacadae decession	
23661 acatcaattg acatgccct ggood	Table 1350 and 1350 a
22721 ctcatattat caccadacty course	
33701 acadtacagt acaagegeag access	The tag sastataatc aggcagagtt
230/1 gratattogg adcoaccage adcasons	cattcagaac ctctqqqatq
33841 gcatattggg aaccaccagt aatatcatc 33901 tcttgtagaa attgaataaa agaaaaatt 33961 caaatgcaat aggttaccgc gctgcgct	cc aacattgtta gttttgaatt agtctgcaaa
33961 caaatgcaat aggttacege getgeget	

34081 34141 34201 34261 34321 34381 34501 34561 34621 34681 34741	tttccatcac gattaaacaa catacaatcc gtataattat gcacaggaga gtccctctaa ggcacacaaa gccctaaact ccgaaactgc ttcctcttc	aagacaagcc cagcaccgaa agacatgtta gcttaatcgt ataaaaaata atacacatac ccacaagctc gacgtaatgg gtcaccaggg tcaccaggg	acagggtctc agttcctcgc gcatcagtta aagtatagca taattatttc aaagcctcat taaagtcact gactaaagtg aaaagtacag aaagtacag	cageteyace ggtgaccage aggagaaaaa aagecacece tetgetgetg cagecatge etceaacete taaaaaatec tttaacttce ttaacttaca	aacaggtgga ctcgtaaaac atgaataagt acagccaaca tcgcggatac tttaggcaac ttaccagaga tccacaatat cgccaaaccc gcaatcccaa acgtcatct cacggcccac tatattattg	cttgatgaag tagcctttga aaagtaaaag gtcgccccg aagtacagcg atatacacaa aacacacacc caagcgtcac cccacggccg acttttaaa
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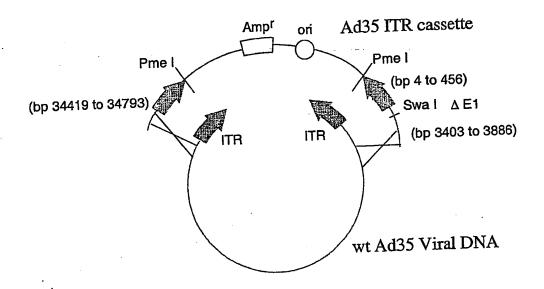


FIG. 3

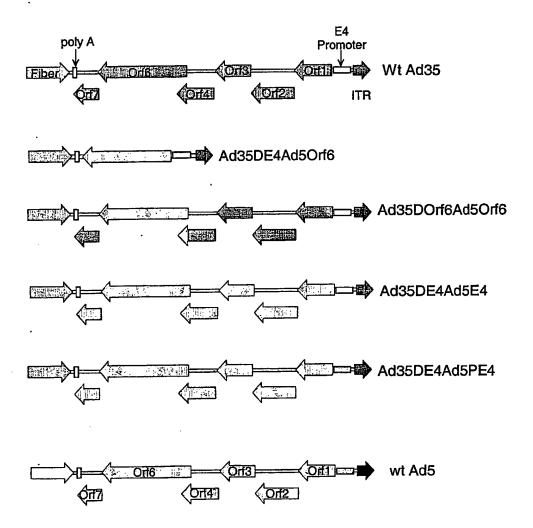


FIG. 4

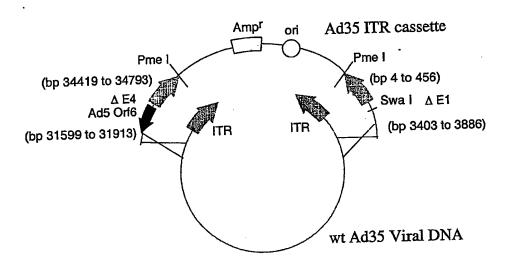


FIG. 5

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121	ttagttcata	occatatat	ggagttccgc	gttacataac	ttacggtaaa	tggcccgcct
191	ggctgaccgc	ccaacgaccc	cccccatto	acgtcaataa	tgacgtatgt	tcccatagta
241	acgccaatag	ggactttcca	ttgacgtcaa	tagatagagt	atttacggta	aactgcccac
301	ttggcagtac	atcaagtgta	tcatatocca	agtacgcccc	ctattgacgt	caatgacggt
361	aaatggcccg	cctggcatta	toccaotac	atgaccttat	gggactttcc	tacttggcag
421	tacatctacg	tattagtcat	coctattacc	atggtgatgc	ggttttggca	gtacatcaat
481	gagagatagat	agcggtttga	ctcacqqqqa	tttccaagtc	tccaccccat	tgacgtcaat
541	gggagtttgt	tttoocacca	aaatcaacgg	gactttccaa	aatgtcgtaa	caactccgcc
601	ccattgacgc	aaatgggggg	taggcgtgta	cggtgggagg	tctatataag	cagagetegt
661	ttagtgaacc	atcagatege	ctggagacgc	catccacgct	gttttgacct	ccatagaaga
721	Caccadascc	gatccagcct	ccacaaccaa	gaacggtgca	ttggaacgcg	gattccccgt
781	accaagagta	agatetaccA	TGGGTGCTAG	GGCTTCTGTG	CTGTCTGGTG	GTGAGCTGGA
841	CAACTGGGAG	AAGATCAGGC	TGAGGCCTGG	TGGCAAGAAG	AAGTACAAGC	TAAAGCACAT
901	TGTGTGGGCC	TCCAGGGAGC	TGGAGAGGTT	TGCTGTGAAC	CCTGGCCTGC	TGGAGACCTC
961	TCAGGGGTGC	AGGCAGATCC	TGGGCCAGCT	CCAGCCCTCC	CTGCAAACAG	GCTCTGAGGA
1021	CCTCACGTCC	CTGTACAACA	CAGTGGCTAC	CCTGTACTGT	GTGCACCAGA	AGATTGATGT
1081	GAAGGACACC	AAGGAGGCCC	TGGAGAAGAT	TGAGGAGGAG	CAGAACAAGT	CCAAGAAGAA
1141	GGCCCAGCAG	GCTGCTGCTG	GCACAGGCAA	CTCCAGCCAG	GTGTCCCAGA	ACTACCCCAT
1201	TGTGCAGAAC	CTCCAGGGCC	AGATGGTGCA	CCAGGCCATC	TCCCCCCGGA	CCCTGAATGC
1261	CTCCCTCAAG	GTGGTGGAGG	AGAAGGCCTT	CTCCCCTGAG	GTGATCCCCA	TGTTCTCTGC
1321	CCTGTCTGAG	GGTGCCACCC	CCCAGGACCT	GAACACCATG	CTGAACACAG	TGGGGGGCCA
1381	TCAGGCTGCC	ATGCAGATGC	TGAAGGAGAC	CATCAATGAG	GAGGCTGCTG	AGTGGGACAG
1441	CCTCCATCCT	GTGCACGCTG	GCCCCATTGC	CCCCGGCCAG	<i>ATGAGGGAGC</i>	CCAGGGGCTC
. 1501	TGACATTGCT	GGCACCACCT	CCACCCTCCA	GGAGCAGATT	GGCTGGATGA	CCAACAACCC
1561	CCCCATCCCT	GTGGGGGAAA	TCTACAAGAG	GTGGATCATC	CTGGGCCTGA	ACAAGATTGT
1621	GAGGATGTAC	TCCCCCACCT	CCATCCTGGA	CATCAGGCAG	GGCCCCAAGG	AGCCCTTCAG
1681	GGACTATGTG	GACAGGTTCT	ACAAGACCCT	' GAGGGCTGAG	CAGGCCTCCC	AGGAGGTGAA
·1741	GAACTGGATG	ACAGAGACCC	TGCTGGTGCA	GAATGCCAAC	CCTGACTGCA	AGACCATCCT
1801	GAAGGCCCTG	GGCCCTGCTG	CCACCCTGGA	GGAGATGATG	ACAGCCTGCC	AGGGGGTGGG
1861	GGGCCCTGGT	CACAAGGCCA	GGGTGCTGGC	TGAGGCCATG	TCCCAGGTGA	CCAACTCCGC
1921	CACCATCATG	<i>ATGCAGAGG</i> G	GCAACTTCAG	GAAÇCAGAGG	AAGACAGTGA	AGTGCTTCAA
1981	CTGTGGCAAG	GTGGGCCACA	TTGCCAAGAA	. CTGTAGGGCC	CCCAGGAAGA	AGGGCTGCTG
2041	GAAGTGTGGC	AAGGAGGGCC	' ACCAGATGAA	GGACTGCAAT	GAGAGGCAGG	CCAACTTCCT
2101	GGGCAAAATC	TGGCCCTCCC	' ACAAGGGCAG	GCCTGGCAAC	TTCCTCCAGT	CCAGGCCTGA
2161	GCCCACAGCC	CCTCCCGAGG	AGTCCTTCAG	GTTTGGGGAG	GAGAAGACCA	CCCCCAGCCA
2221	GAAGCAGGAG	CCCATTGACA	AGGAGCTGTA	CCCCCTGGCC	TCCCTGAGGT	CCCTGTTTGG
2281	CAACGACCCC	TCCTCCCAGI	'AAaataaagc	ccgggcagat	ctgatetge <u>t</u>	grgccrccta
2341	gttgccagcc	atctgttgtt	tgeceeteee	ccgtgccttc	cttgaccctg	gaaggtgcca
2401	ctcccactgt	cctttcctaa	taaaatgagg	aaattgcatc	gcattgtctg	agtaggtgtc
2461	attctattct	ggggggtggg	grggggcagc	acagcaaggg	ggaggattgg	gaagacaata
	gcaggcatgc	tggggatgcg	grgggctcta			
CEO	TD MO. 2					

SEQ ID NO: 2

the base of the safe	aat atgtacattt atattggctc atgtccaaca
1 ccattgcata cgttgtatcc ataccate	act agttattaat agtaatcaat tacggggtca
61 tracegorat grigacatty attacty	act atglacatte dedegger act agttattaat agtaatcaat tacggggtca cgc gttacataac ttacggtaaa tggcccgcct
121 ttagttcata gcccatatat ggagtte	tracgtatot toccatagta
181 ggctgaccgc ccaacgaccc ccgccca	been attracegta aactgcccac
241 acgccaatag ggactttcca ttgacgt	tad tyggsggg ctattgacgt caatgacggt
301 ttggcagtac atcaagtgta tcatatg	-transttat gggactttcc tacttggcag
361 aaatggcccg cctggcatta tgcccag	at atgetatac gattttagca gtacatcaat
421 tacatctacg tattagtcat cyctate	btterporte tecaceccat tgacgtcaat
481 gggcgtggat agcggtttga ctcacgg	gga coossaga aatgtcgtaa caactccgcc
541 gggagtttgt tttggcacca adattad	tratataag cagagetegt
601 ccattgacgc aaatgggegg taggege	gta oggogget gttttgacct ccatagaaga
661 ttagtgaacc gtcagatcgc ctggaga	ttgge category ttggaacaca gattccccgt
721 caccgggacc gatccagcet ecgegge	COMPANY CONTROL CONTRO
781 gccaagagtg agatcgatct aagtaag	TO THE TOTAL CONTROL CONTROL COGGACTTCT
841 GCCTGAGGCT ACAGCTCTCC CIGGGC	COCCCAAGA GCTGCAGCCT GCACAGACAG
901 GGAACCGCGA GGCAGCCGAG GCCCTGG	TOTAL AMERICAN MICCO COTOTOTACO GTGACAGCTG
961 CCGCCAAGAA CCTCATCATC TTCCTGC	THE ARCTICECCC TEAGATACCC CTGGCCATGG
1021 CCAGGATCCT AAAAGGGCAG AAGAAGG	ACA CARACA ACCO ACACAAACAT GTGCCAGACA
1081 ACCGCTTCCC ATATGTGGCT CTGTCCA	THE COMMENT OF CANCETY CAG ACCATTGGCT
1141 GTGGAGCCAC AGCCACGGCC TACCTG	TGCG GGGTCATCTCCG CGCCAACGAG GTCATCTCCG
1201 TGAGTGCAGC CGCCCGCTTT AACCAG	TOCK HOLLOWING GETAACCACC ACACGAGTGC
1261 TGATGAATCG GGCCAAGAAA GCAGGG	ARGI CHOTOGOTONA CCCCAACTGG TACTCGGACG
1321 AGCACGCCTC GCCAGCCGGC ACCTAC	CATCCCTACG CAGCTCATCT
1381 CCGACGTGCC TGCCTCCGCC CGCCAG	GAGG GGIGCCCAAA GTACATGTTT CGCATGGGAA
1441 CCAACATGGA CATTGACGTG ATCCTA	THE GAGGEGARGE CACCAGGCTG GACGGGAAGA
1501 CCCCAGACCC TGAGTACCCA GATGAC	TACA SCOTTCCCC CTATCTCTGG AACCGCACTG
1561 ATCTGGTGCA GGAATGGCTG GCGAAG	TOTAL MARCECATCT CATCCCTCTC TTTGAGCCTG
1621 AGCTCATGCA GGCTTCCCTG GACCCG	TOTAL TONCOUNTER CCCCTCCCTG ATGGAGATGA
1681 GAGACATGAA ATACGAGATC CACCGA	ACCOCCCCC CTTCTTCCTC TTCGTGGAGG
1741 CAGAGGCTGC CCTGCGCCTG CTGAGC	AGGA ACCCCCTTA CCGGGCACTG ACTGAGACGA
1801 GTGGTCGCAT CGACCATGGT CATCAL	GARA GCHOCOTTAC CAGCGAGGAG GACACGCTGA
1861 TCATGTTCGA CGACGCCATT GAGAGG	TOTAL MORE TITLES AGECTACCC CTGCGAGGGA
1921 GCCTCGTCAC TGCCGACCAC TCCCAC	THE COCCOCACAG GAAGGCCTAC ACGGTCCTCC
1981 GCTCCATCTT CGGGCTGGCC CCTGGC	ARGO COCCOCCC CCCCCCCGAT GTTACCGAGA
2041 TATACGGAAA CGGTCCAGGC TATGTC	TOTAL AGGREGACIANT COCOUNTEGAC GAAGAGACCC
2101 GCGAGAGCGG GAGCCCCGAG TATCGG	CAGE AGTERICACE GEOGRACETG GTTCACGGCG
2161 ACGCAGGCGA GGACGTGGCG GTGTTC	TOTAL MEGGETTECC CCCCTGCCTG GAGCCCTACA
2221 TGCAGGAGCA GACCTTCATA GCGCAC	TOTAL TOUCCULOUS CONCENTRACECE
2281 CCGCCTGCGA CCTGGCGCCC CCCGCC	grantagg tggcccccc tgaattggaa
2341 tggtccccgc gttgcttcct ctgct	thetaethee caecatete ttetteece
2401 togatoagaa ttgatotgat ougos	transtace actateett cetaataaaa
2461 ctccccgtg ccttccttga ccctg	gaagg tgetaetet attctggggg gtggggtggg
2521 tgaggaaatt gcatcgcatt gtctg	gaagg tgctactccc actgsservices agtag gtggggtggg gtggggtggg gaaga caatagcagg catgctgggg atgcggtggg
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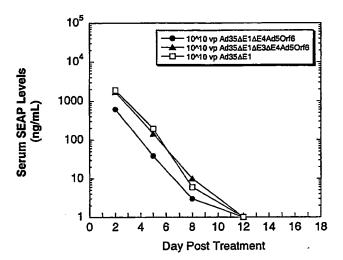


FIG. 8

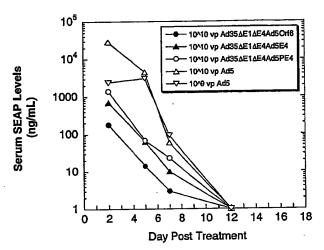


FIG. 9

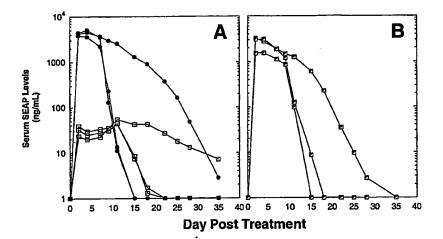


FIG. 10A-B

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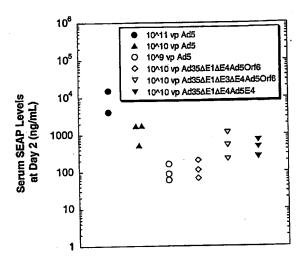


FIG. 11

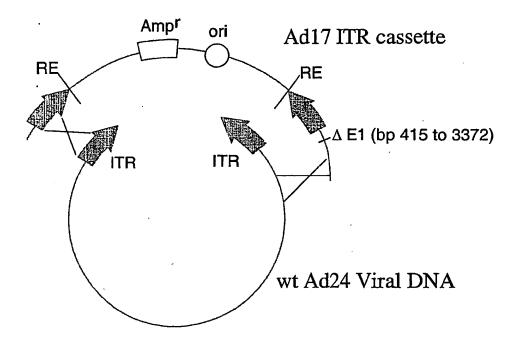


FIG. 12

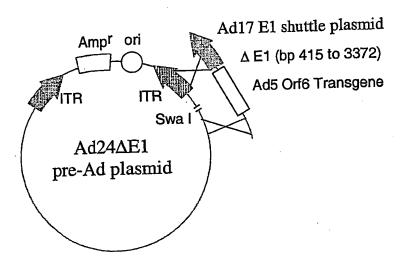


FIG. 13

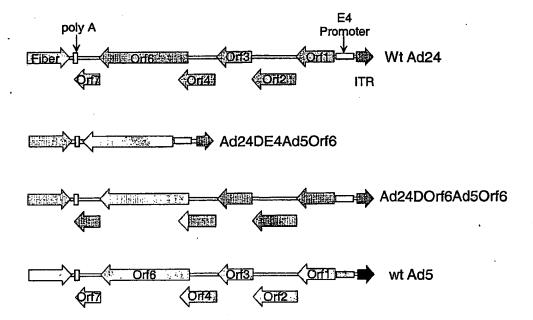


FIG. 14

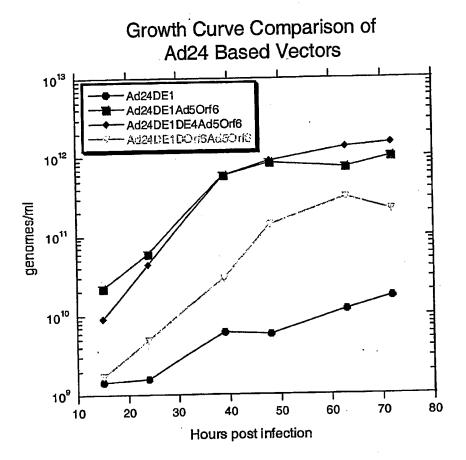


FIG. 15

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121	acggctaacg	atcaccacaa	aggcataacc	tagcccggaa	gcaagtcgcg	gggctgatga
101	cgtataaaaa	accoracttt	aggogogge	accocccatt	ttcccacaac	cacqcccqqa
T 8 T	tatgaggtaa	ttataaaaaa	atacaaataa	aattacctca	ttttaacaca	aaaactgaat
241	gaggaagtga	anagtgaga	ataccontcc	cacccadade	ggaatattta	ccaaaaacca
301	agagactttg	adaytyddad	tagaaattta	cattacaata	tttttcaca	aatttcccc
361	tccgtgtcaa	accyattacy	ttatataaca	gattgcggcg	tccacagggt	atttaaacca
421	tccgtgtcaa	agteeggtgt	cratteres	gaccagecga	agagget	ctgagetecg
481	gtcgagcccg ctcccagagt	tcaagaggcc	actettgagt	gccagcgagt	tetteaacta	tacctattaa
541	ctcccagagt	ctgagaaaaa	Lyayacacct	gogooccocc	ttagaggagg	aactgcatcc
601	catggccgca	ttattgetgg	aggarrargr	gagtatatat	ceggaggaeg	tagatgccca
661	atctccattt	gagetgggae	etacaettea	gyacctatat	gatttggagg	trattcttca
721	tgatgacgac	ccgaacgaag	aggetgtgaa	LLLaatatt	ccayaacctc	tatcacccat
781	ggctgacata	gccagcgaag	etgtacetae	accaccicat	tataaaaaa	attttcctcc
841	acctgaattg	gaagaggagg	acgagctaga	ceteegatgt	catgaggaag	aatatactta
901	cagcgattca	gaggacgaac	agggcgagca	gageauggeu	ctaatettaa	aacacyctcy
961	tgtggttgtg	gaagagcatt	ttgtgttgga	caateetgag	gegeeeggge	atatatatta
1021	atcctgccag	taccaccggg	ataagaccgg	agacacgaac	geeteetgeg	coccycycta
1081	catgaaaaag	aacttcagct	ttatttacag	taagtggagt	gaatgtgaga	gagactgagt
1141	gcttaacaca	taactgggta	atgcttaaac	agctgtgcta	agtgtggttt	atttttgtt
1201	ctaggtccgg	tgtcagagga	tgagtcatca	ccctcagaag	aagaccaccc	gtgtcccct
1261	gagetgtcag	acaaacacc	cctgcaagtg	cacagaccca	ccccagtcag	acccagtggc
1321	gagagggag	cagctgttga	aaaaattgag	gacttgttac	atgacatggg	tggggatgaa
1381	cctttggacc	tgagcttgaa	acccccagg	aactaggctc	agctgtgctt	agtcatgtgt
1441	aaataaagtt	otacaataaa	agtatatgtg	acgcatgcaa	ggtgtggttt	atgactcatg
1501	gacataactt	agtcctatat	aagtggcaac	acctgggcac	tggggcacag	accttcaggg
1561	agttcctgat	ggatgtgtgg	actatccttg	cagactttag	caagacacgc	cggcttgtag
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1681	gtctggtgta	cacagttaag	aaggattata	acgaggaatt	tgaaaatctt	tttgctgatt
1741	actetagest	octagattct	ctaaatctcg	gccaccagtc	ccttttccag	gaaagggtac
1201	tecacageet	tgatttttca	agcccagggc	gcactacagc	cggggttgct	tttgtggttt
1861	ttctggttga	caaatggagc	cagaacaccc	aactgagcag	gggctacatt	ctggacttcg
1921	cagocatgoa	cctgtggagg	gcatgggtga	ggcagcgggg	acagagaatc	ttgaactact
1981	ggcttataca	gccagcagct	ccaaatcttc	ttcgtctaca	cagacaaaca	tccatgttgg
2041	accaacaaat	gaggcaggcc	atggacgaga	acccgaggag	cggcctggac	cctccgtcgg
2101	aagaggagct	ggattgaatc	aggtatccag	cctgtaccca	gagettagea	gggtgctgac
2161	atccatggcc	aggggagtga	agagggagag	gagcgatggg	ggcaataccg	ggatgatgac
2221	cgagctgacg	accaacctaa	tgaatcgcaa	gcgtccagag	cgcattacct	ggcacgagct
2221	acagatggag	tatagggata	aggtgggct	gatgcaggat	aaatatggcc	tggagcagat
2201	aaaaacccac	taattaaacc	cagatgagga	ttgggaggag	gccattaaga	aatatgccaa
2/101	gatagccctg	caccaaatt	acaaatacaa	ggtgaccaag	acqqtqaata	tcagacatgc
2401	ctgctacatc	terrareaca	addcadaddt.	ggtcatcgat	accctggaca	aggccgcctt
2401	caggtgttgc	atratorna	taaaaaccaa	agtgatgaat	atgaattcca	tgattttcat
2521	gaacatgaag	ttoastaaa	araarttaa	tagagatgata	ttcatggcca	acagtcacat
720T	gaccetgeac	cactaceatt	tetteggett	caacaatato	tacacagaga	tetagagege
2041	tgctaagatc	ggctgcagtt	actttataa	ctactacata	aacataatca	daadacccaa
2/UI	gagcgagatg	aggggatgta	agteteatgy	traraaatrr	tacctoooad	tetetacega
2761	gagegagatg	tetgtgaage	agigigigic	cgagaaacgc	caccegggag	acctaatas
2821	gggcaatgct	agagtgagac	attyctctc	cccgyagacg	accostosco	gcctggcgaa
2881	gggcacagcc	tetetgaage	ataatatggt	gaagggerge	acggacgage	traceteres
2941	catgctgaca	tgcgactcgg	gggtetycea	taccetgaay	adcatttatg	reateracet
3001	ccccggaag	aagtggccag	tgtttgagaa	taacctactg	accaagigee	acatycacct
3061	gggcgccaga	aggggcacct	tccagccgta	ccagtgcaac	tttageeaga	ccaagetget
3121	gctggagaac	gatgccttct	ccagggtgaa	cctgaacggc	accurraca	Lggatgtete
3181	ggtgtacaag	atcctgagat	acgatgagac	caagtccagg	gtgcgcgctt	gcgagtgcgg
3241	gggcagacac	accaggatgc	aaccagtggc	cctggatgtg	accgaggagc	rgaggcccga
3301	ccacctggtg	atggcttgta	ccgggaccga	gttcagctcc	agtggggagg	acacagatta
3361	gaggtaggtt	gagtattagt	gggcgtggct	aaggtgacta	taaaggcggg	tgtcttacga
3421	agatetttt	gcttttctgc	agacatcatg	aacgggactg	gcggggcctt	cgaagggggg
3481	ctttttagcc	cttatttgac	aacccgcctg	ccgggatggg	ccggagttcg	tcagaatgtg
3541	atoggatcga	caataaacgg	gcgtccagtg	cttccagcaa	attcctcgac	catgacctac
3601	gcgaccgtgg	ggaactcgtc	gctcgacagc	accgccgcag	ccgcggcagc	cgcagccgcc

FIG. 16A-1

2661		cgagactggc	ttaaaaataa	atorccagca	gcagcagtag	cccctctata
3001	atgacagcya	cyayactyyc		ataccetae	taaccaaact	ggaaggggtg
3721	cccagttcca	tcatcgccga	ggagaaaccg		taggeegagee	ggaagooos
3781	agccgccagc	tggccgccct	gacccagcag	gratecyaye	Leegegaaca	geageageag
3841	caaaataaat	gattcaataa	acacagattc	tgattcaaac	agcaaagcac	Cittatiati
3001	tattttttca	cacacaataa	accetaatee	acctctcccg	accattgaga	gracaaraa
2061	++++++ccaa	cacccontag	aggtgggatt	ggatgttgag	gtacatgggc	atgageeegt
4021	cecanagata	gaggtaggag	cactgcatgg	cctcgtgctc	tggggtcgtg	tigiagaiga
4001	tocactcata	acsadaacac	tagacataat	actagatgat	gtccttgagg	aggagactya
4141	tagagagaga	gagccccttg	atatagatat	taacaaaaca	gttgagctgg	gagggatgca
4741	tacaaaaaaa	gatgatgtgg	agtttggcct	ggatcttgag	gttggcgatg	ttgccaccca
420I	cgcgggggga	ggggttcatg	ttatacada	ccaccagaac	gatatagece	gtgcacttgg
4201	gateeegeet	atgcaacttg	gegeagga	catasaaaa	tttggagacg	cccttatacc
4321	ggaacttgtc	ttccatgcac	gaayyyaary	tastaacast	gaaccataa	actacaactt
4381	cacccaggtt	ttecatgeac	teaccatga	cataattata	ctcctacata	agatcatcat
4441	tggcaaagac	gtttctgggg	tcayayacat	terangetta	aaaaaaata	atteceteaa
4501	aagacatttt	aatgaatttg	gggcggaggg	tgecagattg	ggggacaatg	tagaaaaaa
4561	gccccggggc	gaagttcccc	tcacatattt	gcatctccca	ggettteate	ccggaggggg
4621	ggatcatgtc	cacctgcggg	gcgatgaaaa	aaacggtttc	cggggcgggg	gcgacgagec
4681	acasaasaa	caggtttctc	aacagctggg	acttgccgca	cccggtcggg	ccgtagatya
1711	ccccgatgac	gggttgcagg	taataattca	aggacatgca	gctgccgtcg	teeeggagga
4801	agggggggggg	ctcgttgage	atotetetoa	cttggaggtt	ttcccggacg	agctcgccga
4861	ggagggggtc	ccccccacc	gagagcagct	cttgcaggga	agcaaagttt	ttcaggggct
4921	tragrecente	agceatagge	atcttqqcqa	gggtctgcga	gaggagttcg	aggeggteee
4001	agageegee	gacgtgctct	acoccatctc	gatccagcag	acttcctcgt	ttcgggggtt
E041	agagecegge	cgactgtagg	acacaaaaca	ataggcatcc	agcgctgcca	gcgtcatgtc
5101	gggacgactg	ctcagtgtcc	acatagagaeg	gatetecate	accottoaago	gatagacccc
2101	cuccayyyu	cttgcaaggg	tacacttaaa	actcatcctq	ctggtgctga	aacgggcacg
2101	gggetgtgeg	tgcgcgtcgg	casastages	attacceta	acctcotagt	tgagggcctc
5221	gtcttcgccc	tgegegtegg	cyagatagea	attaceaca	caccacaca	caaaacaaaa
5281	ggcggcgtgg	cccttggcgc	ggagettgee	cccggaagag	caccacaca	cdaaadcatc
5341	gagggattgc	agggcgtaga	gerraggrae	gagaaagacg	gacteggggg	caaaatactc.
5401	cgctccgcag	tgggcgcaga	"cggtctcgca	etegaceage	caygrager	coggetgete
5461	ggggtcaaaa	accagttttc	ccccgttctt	tttgatgcgc	ttettaeete	gegteteat
5521	gagtetgtgt	ccacactcaa	tgacaaacag	gctgtctgtg	tccccgtaga	cggacttgat
5581	gaacctatee	tacagagaca	tecegeggte	ctcctcgtag	agaaactcgg	accactetga
56/1	dacdaaddcd	cocotccaco	ccaagacaaa	ggaggccacg	tgcgaggggt	ageggtegtt
5701	gtccaccagg	gggtccacct	tttccacggt	atgcagacac	atgtccccct	cctccgcatc
5761	caagaaggtg	attggcttgt	aggtgtaggc	cacgtgaccc	ggggtccccg	acgggggggc
5921	ataaaaaaaa	acaaatetat	actcatcctc	actctcttcc	gcgtcgctgt	ccacgagcgc
5001	cacctattaa	ggtaggtatt	ccctttcgag	agcgggcatg	acctcggcac	tcaggttgtc
5001	caycogcogg	aacgaggagg	atttgatgtt	aacttaccct	geogeaatge	tttttaggag
234T	agttttctaga	atctggtcag	assagactat	ttttttattg	tcaagcttgg	togcgaagga
POOT	actitication	gcgttggaga	gaagettggc	gatggatctc	atggtctgat	ttttgtcacg
909T	gecatagagg	tecttggeeg	gaageeegge	ctonacatac	tenenenena	cocacttcca
6121	geoggetege	teettggeeg	cyatyttyay	ceggacacac	scacaccaac	cacaattata
6181	ttcggggaag	acggtggtgc	getegteggg	cacyaccecy	acgegeeage	taatccaaca
6241	cagggtgacc	agatccacgc	tggtggccac	etegeegege	aggggcccgc	cotcagea
6301	gaggcgtccg	cccttgcgcg	agcagaacgg	gggcagcaca	tcaagcagat	gcccgccagg
6361	ggggtccgca	tcgatggtga	agatgcccgg	acagagttcc	ttgtcaaaat	aatcgatttt
6421	tgaggatgca	tcatccaagg	ccatctgcca	ctcgcgggcg	gccagcgctc	gctcgtaggg
6481	attaagagag	ggaccccagg	gcatgggatg	cgtcagggcg	gaggcgtaca	Lgccgcagat
6541	gt.cgtagaca	tagatgggct	ccgagaggat	gccgatgtag	gtgggataac	agegeeeee
6601	acagatacta	gegegeaegt	agtcatacaa	ctcgtgcgag	ggggccaaga	aggcggggcc
6661	gagattggtg	cgctggggct	actcaacaca	gaagacgatc	tggcgaaaga	tggcatgcga
6721	attagaggag	atggtgggcc	gttggaagat	gttaaagtgg	gcatgaggca	gacgaaccga
6701	gttggaggag	aagtgcgcgt	aggagtettg	cagettggeg	acgagetegg	cggtgacgag
60/01	geogeggatg	gcgcagtagt	CCSGCGtttc	gcggatgatg	tcataaccco	cctctccttt
0841	gacytccaty	agctcgcggt	traccrosts	ctcctcatce	teetteeaet	actcccaaa
6901	CTTCCCCCAT	agulugugu	-uayyycyta	. cccccacata	tanasatnot	tcacqqcctt
6961	cgggaatcct	cgatcgtccg	cacggcaaga	, gcccagcatg	tagaaacygt	tacaagacaa
7021	gtagggacag	cagecettet	ccacggggag	ggcgcaagct	gagcygcct	. cycygaycya
7081	ggtgtgcgtc	agggcgaagg	tatccctgac	catgactttc	aayaactygt	tacceydaalC
7141	. cgagtcgtcg	cagccgccgt	gctcccagag	ctcgaaatcg	gracacrer	. ccgagagggg
7201	gttaggcaga	. gcgaaagtga	cgtcattgaa	ı gagaatettg	r cctgcccgcg	gcatgaaatt
7261	gcgggtgatg	cggaaagggc	ccgggacgga	, ggctcggttg	, ttgatgacct	. gggcggcgag
						•

7221	ascastatea	tcgaagccgt	taatattata	cccgacgatg	tagagttcca	tgaatcgcgg
7361	gacgaccccg	atgtgcggca	actttttaa	ctcctcgtag	gtgaggtcct	cggggcaatg
/38I	geggeettta	tgctcgagcg	gettettgag	gagatgaga	ttggcttgca	tgaatgaagc
7441	cagteegtge	cgggccataa	~~~totaccccg	ctcatcacas	aadaddcdda	actoctoocc
7501	ccagageteg	cgggccacaa	gggtetggag	caccaccac	agateceact	cccagcgatc
7561	cacggccatc	ttttctgggg	cyacycayca	gadag caagg	aggtotogot	cccccagaa
7621	ccagcgtaag	cgcacggcta	gategegage	gagggcgacc	gecoegge	aggtgtaggt
7681	tttcataacc	agcataaagg	ggacgagctg	cttgccgaag	yaccccatcc	ttaaaaaaa
7741	ttctacatcg	taggtgacaa	agageegete	cgtgcgagga	tyayayeeya	ctgggaagaa
7801	ctggatttcc	tgccaccagt	tggacgagtg	gctgttgatg	tgatgaaagt	agaaateetg
7861	ccaacaaacc	gaggagtcgt	actgatgctt	gtaaaagcgt	ccgcagtact	egeagegeeg
7921	cacgggctgt	acctcatcca	cgagatacac	agcgcgtccc.	ttgaggagga	actteaggag
7981	taacaaccet	aactaataat	tttcatgttc	gcctgcgtgg	gactcaccct	ggggctcctc
RN41	daddacadad	aggetgaega	acccacacaa	gagccaggtc	cagatetegg	cgcggcgggg
R101	acadagaaca	aagacgaggg	cacacaatta	ggagctgtcc	atggtgtcgc	ggagatecay
9161	atccagaaac	agggttctga	aattaacctc	gtagaggcgg	gtgagggcgt	gcttgagatg
8221	cadatogtac	ttgatctcca	caaataaatt	ggtggctgtg	tccacgcatt	gcatgageee
8281	ateactacac	ggggccacga	ccataccaca	gtgcgctttt	agaagcggtg	cegeggaege
23/11	acteceases	acaacaacaa	ttccaacccc	gcgggcaggg	gcggcagagg	cacgreggeg
8401	tagegetegg	gcaggtcccg	atactacacc	ctgagagcgc	tggcgtgcgc	gacgacgcgg
8461	conttoacat	cctggatctg	ccacctctgc	gtgaagacca	ccggccccgt	gactttgaac
8521	ctgaaagaca	gttcaacaga	atcaatctcg	gcgtcattga	cggcggcctg	acgcaggate
8581	tettacacat	cgcccgagtt	atcetagtag	gcgatctcgg	acatgaactg	ctcgatctcc
96/1	tectectors	gatcgccgcg	acccacacac	tccacggtgg	cggcgaggtc	attggagatg
0701	ccaccatga	gctgcgagaa	aacacccaaa	ccactctcat	tccagacgcg	gctgtagacc
0761	aggtagagat	cggcgtcgcg	cacacacata	accacctgcg	cgaggttgag	ctccacgtgc
0/01	acgeeeege	cggcgtagtt	acacaacac	tagaagaggt	agtttagggt	ggtggcgatg
0041	tactogatas	cgaagaagta	catdatccad	caacacaaaa	gcatctcgct	gatgtcgccg
0001	tyctcygtga	gcctttccat	aacctcataa	aaatccacag	cgaagttgaa	aaactgggcg
0941	atggeeteea	agaccgtgag	ctcatcatca	aggagggtga	tgagttcggc	gatggtggcg
9001	ttgegggeeg	gctcgaaatc	cccaccaccc	tectectett	cctcttcttc	catgacgacc
9001	egeaectege	tttcttcctc	tacagggggc	aataataaca	aaacccaaca	acqacqqcqa
9121	CCCCCCCC	gacggtcgac	caaccctca	atcatctccc	cacaacaaca	acqcatqqtt
9101	taaataaaa	cgcgaccccg	ttccccacca	cacaacataa	agacgccgcc	ggtcatctcc
9241	ceggigacyg	gegggteece	attanaceae	dadaddddcac	tgacgatgca	tcttatcaat
9301	cygraarygy	gggacgtgag	cacatcasas	tcgaccggat	cogagaatet	ttcgaggaaa
9361	tgcggtgtag	aatcgcagtc	cgcgccgaga	ctcaaacaca	taggagggggt	gtggacgctg
9421	gegtetagee	ggttgctgat	geauggeaug	aagtaggggt	ttttaaggcg	acagatagta
9481	ttagaattge	ccaggtcctt	gatgtaattg	tactaggtgc	gaagegete	ggccatgccc
9541	gcgaggagga	cctgacaccg	gggtcccgcc	ttataataat	catocatoao	cctctcaatg
9601	caggcctggc	cetgacaeeg	gettaggett	cagatasacc	caegaacgag	gagcggctgc
9661	tcatcactgg	cggaggcgga	greceedatg	cgggcgaccc	cctattacec	acaaataaaa
9721	acgagcgcca	ggtcggcgac	gacgegeteg	gcyagyacyg	congregate	gegggegagg
9781	gtgtcctgga	agtcgtccat	gccgacgaag	cggtggtagg	ccccggcgcc	gacggcgcag
9841	gtgcagttgg	ccatgagcga	ccagttgacg	grergeagge	cgggccgcac	gacccccgag
9901	tacctgagcc	gcgagaaggc	gcgcgagtcg	aagacacagc	cgttgcaygt	gegeacgagg
9961	tactggtatc	caactaggaa	gtgcggcggc	ggerggeggr	agageggeea	gegetgggtg
10021	gccggcgcgc	ccggggccag	greeregage	atgaggeggt	ggtageegta	gaggtagtgg
10081	gacatccagg	tgatgccggc	gacaaraara	gaggegegeg	ggaactcgcg	gacgcggttc
10141	cagatgttgc	gcagcggcag	gaaatagtcc	atggtcggca	eggtetggee	gytgagacgc
10201	gcgcagtcat	tgacgctcta	gaggcaaaaa	cgaaagcggt	tgageggget	etteeteegt
10261	agcctggcgg	aacgcaaacg	ggttaggccg	cgtgtgtacc	ccggttcgag	teeeetegaa
10321	tcaggctgga	gccgcgacta	acgtggtatt	ggcactcccg	tetegaceeg	agcccgatag
10381	ccaccaggat	acggcggaga	geeetttttg	ccgaccgagg	ggagtcgcta	gacttgaaag
10441	списсивава	ccccaccaaa	tagtggctcg	cgcccgtagt	ctggagaagc	tttgccaggg
1.0501	ttgagtcgcg	gcagaacccg	gttcgcggac	ggccgcggcg	agcgggactt	ggtcaccccg
10561	ccgatttaaa	gacccacage	cagccgactt	ctccagttac	gggagcgagc	CCCCCCCCCCC
10621	ctttttgcca	gatgcatccc	qtcctgcgcc	aaatgcgtcc	CACCCCCCC	ccggcgacca
10681	ccgcgaccgc	gaccataaca	ggcgccggcg	ctgtagcccc	gccacagcag	acagagatgg
10741	acttggaaga	gggcgaaggg	ctggcgagac	tgggggcgcc	gtccccggag	cgacaccccc
10801	acatacaact	gcagaaggac	gtgcgcccgg	cgtacgtgcc	tgcgcagaac	ctgttcaggg
10861	accocaocoo	GGAGGAGCCC	gaggagatgc	gcgactgccg	ttttcgggcg	ggcagggagc
10921	tacacaagaa	cctggaccgc	cagcgcgtgc	tgcgcgacga	ggatttcgag	ccgaacgagc

FIG. 16A-3

		cagccccgcg	racacacaca	taacaacaac	caacctggtg a	acggcctacg
11041	agcagacggt	cgaggaggtg	cgcaaccccc	tgatgcacct	gtgggacctg	gcggaggcca
11101	taatcgcgcg	cccggacagc	nogatatan	caacacaact	attectaata	gtgcagcaca
11161	tcgtgcagaa	cccggacage	aageetetga	tactaaacat	caccagaccc	gagggccgct
11221	gcagggacaa	gctgatcaac	agggaggcgc	gcatcgtagt	gcaggagcgc	agcctgagcc
11281	ggctgctgga	gctgatcaac	attitudaga	cartactasa	cctgggcaag	ttttacgcgc
11341	tggccgagaa	ggtggcggct	attaattatt	tagagagaga	ggtgaagata	gacagetttt
		~~~~~~~~~~	actagacalu	accucacaca	acacggaace	
			- Madeconeco	COLLUCION	90909900	3
1440	1 cagaatgca	a ccagggctgc g ctgaaaagga	tatastast	t agaggagae	a cacaagatga	a aagtaagaga
1446	l gaacaagcg	g ctgaaaagga g tcatagatgg	a cocygreat	a accetetac	c gaagttggt	a cctgtcctat
1452	1 agctataat	g tcatagatgo g accccgagaa	acceptange	r teatagaea	c toctcacca	cccggacgtc
1458	1 acctacggg	g acceegagaa	ı gygygtyca	g cogoggaes		

FIG. 16A-4

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14641	acctgcggcg	cogagcaagt	ctactggtcg	ctgccggacc	tcatgcaaga	ccccgtcacc
14701	ttcccctcta	cccagcaagt	caccaactac	cccataatta	gcgccgagct	catgcccttc
14761	cacaccasaa	actttacaa	cgacctcgcc	gtctactccc	agctcatccg	cagctacacc
1/1921	teceteacce	acatetteaa	ccacttccc	gacaaccaga	tcctctgccg	tecaceaca
1/001	ccccccatce	ccacaatcaa	tgaaaacgtg	cctactctca	cagatcacgg	gacgctaccg
14041	ctcaccacca	atatoggaag	agtccacca	ataaccatca	ctgacgcccg	teaccacace
16001	teteestees	tatagagaga	agtecagega	atcacacaa	gcgtgctttc	carteresec
T200T	tgteectacg	tetacaayye	cctgggcata	accastascs	ccaactaaaa	tottactacc
12001	ttctaaaaaa	tgtetattet	catcucyccc	agcaacaaca	ccggctgggg	coccactagg
					agcacccegt	
12181	ggccacttcc	gegeteeetg	gggegettae	aagegeggge	ggacttctac	cyccyccycy
15241	cgcaccaccg	tegaegaegt	categaeteg	gradicacca	acgcgcgcaa	ccacaccccc
15301	gccccctcca	eegtggaege	ggtcatcgac	agegraggragg	ccgacgcgcg	cgactatgee
15361	agacgcaaga	geeggeggeg	aeggategee	aggegeeace	ggagtacgcc	cyccacycyc
15421	gccgcccggg	etetgetgeg	cegegeeaga	egeaegggee	gccgggccat	gatgegagee
15481	gcgcgcgcg	cegecaetge	acceccegca	ggeaggaete	gcagacgagc	ggccgccgcc
15541	gctgccgcgg	ccatttctag	catgaccaga	eccaggegeg	gaaacgtgta	etgggtgtgt
15601	gactccgtca	cgggcgcgcg	egrgeeegrg	egeaceegre	ctcctcgtcc	cugatutaat
15661	gcttgtgtcc	tececegeaa	gegaegaege	caaagegeaa	aatcaaggag	gagatgetee
15721	aggtcgtcgc	cccggagatt	tacggaccac	eccaggegga	ccagaaaccc	cycaaaatca
15781	agcgggttaa	aaaaaaggat	gaggtggacg	agggggcagt	agagtttgtg	egegagiteg
15841	ctccgcggcg	gcgcgtaaat	tggaaggggc	gcagggtgca	gcgcgtgttg	eggeceggea
15901	cggcggtggt	gtttacgccc	ggcgagcggt	cctcggtcag	gagcaagcgt	agetatgaeg
15961	aggtgtacgg	cgacgacgac	atcctggacc	aggcggcgga	gcgggcgggc	gagttegeet
16021	acgggaagcg	gtcgcgcgaa	gaggagctga	tetegttgee	gctggacgag	agcaacccca
16081	cgcctagcct	gaagcccgtg	accetgeage	aggtgctgcc	ccaagcagtg	ctgctgccga
16141	gccgcggggt	caagcgcgag	ggcgagaata	tgtacccgac	catgcagatc	atggtgccca
16201	agegeeggeg	cgtggaagaa	gtgctggaca	ccgtgaaaat	ggatgtggag	cccgaggtca
16261	aggtgcgccc	catcaagcag	gtggcgccgg	gccrgggcgr	gcagaccgtg	gacattcaga
16321	tccccaccga	catggatgtt	gacaaaaaac	cctcgaccag	catcgaggtg	cagaccgacc
16381	cctggctccc	agcctccacc	getgeegtet	ccacttctac	cgccgccacg	gctaccgagc
16441	ctcccagaag	gcgaagatgg	ggccctgcca	accggctgat	gcccaactac	gtattgcatc
16501	cttccattat	cccgacgccg	ggctatcgcg	gcacccggta	ctacgccagc	cgcaggcgcc
16561	cagccagcaa	acgccgccgc	cgcaccgcca	cccgccgccg	tctggcccc	gcccgcgtgc
16621	gccgcgtaac	cacgcgccgg	ggccgctcgc	tegttetgee	caccgtgcgc	taccacccca
16681	gcatccttta	atccgtgtgc	tgtgatactg	ttgcagagag	atggctctca	cttgccgcct
16741	gcgcatcccc	gtcccgaatt	accgaggaag	atcccgccgc	aggagaggca	tggcaggcag
16801	cggcctcaac	cgccgccggc	ggcgggccat	gcgcaggcgc	ctgagtggcg	getttetgee
16861	cgcgctcatc	cccataatcg	cggcggccat	cggcacgatc	ccgggcatag	cttccgttgc
16921	gctgcaggcg	tcgcagcgcc	gttgatgtgc	gaataaagcc	tctttagact	ctgacacacc
16981	tggtcctgta	tatttttaga	atggaagaca	tcaattttgc	gtccctggct	ccgcggcacg
17041	gcacgcggcc	gttcatgggc	acctggaacg	agatcggcac	cagccagctg	aacgggggcg
17101	ccttcaattg	gagcagtgtc	tggageggge	ttaaaaattt	cggctcgacg	ctccggacct
					aagggaaaag	
17221	agaacttcca	gcagaaggtg	gtggacggcc	tagcctcggg	cattaacggg	gtggtggaca
17281	tagcaaacca	ggccgtgcag	cgcgagataa	acageegeet	ggacccgcgg	ccgcccacgg
17341	tggtggagat	ggaagatgca	actcctccgc	cgcccaaggg	cgagaagcgg	ccgcggcccg
17401	acgcggagga	gacgatcctg	caggtggacg	agccgccctc	gtacgaggag	gccgtcaagg
17461	ccggcatgcc	caccacgcgt	atcatcgcgc	cactggccac	tggtgtaatg	aaacccgcca
17521	cccttgacct	gcctccgcca	cccacgcccg	ctccaccgaa	ggcagctccg	gttgtgcagc
17581	cccctcctgt	ggcgaccgcc	gtgcgccgcg	tccccgcccg	ccgccaggcc	cagaactggc
17641	agagcacgct	gcacagtatc	gtgggcctgg	gagtgaaaag	tctgaagcgc	cgccgatgct
17701	attgagagag	aggaaagagg	acactaaagg	gagagcttaa	cttgtatgtg	ccttaccgcc
17761	agagaacgcg	cgaagatggc	taccccctcg	atgatgccgc	agtgggcgta	catgcacatc
17821	gccgggcagg	acgcctcgga	gtacctgagc	ccgggtctgg	tgcagtttgc	ccgcgccacc
					cggtggctcc	
17941	gtgaccacgg	accggtccca	gcgtctgacg	ctgcgctttg	tgcccgtgga	tcgcgaggac
18001	accacgtact	cgtacaaggc	gcgcttcact	ctggccgtgg	gcgacaaccg	ggtgctagac
18061	atggccagca	cttactttga	catccgcggc	gtcctggacc	gcggtcccag	cttcaaaccc
18121	tactcgggca	cggcttacaa	cagcctggcc	cccaaaggcg	ccccaactc	tagtcagtgg
18181	gaacaagcta	aagctaccaa	tgccggtcaa	aaggaaactc	acacatttgg	agtagccgct
18241	atgggcggag	aagacattac	agtgaaaggt	cttcaaattg	gaactgatga	aactaaggaa

FIG. 16A-5

40204		atgaaattt	Focadatcaa	acattccagc	cagaacctca a	agtgggagaa
18421	atgaagccat	gttatggete	teacgegaga	assatteete	atattacaat (	gatttcttt
18481	tttacacttg	atgaaaaagg	teagecaace	addatteetg	caratattot (	catgtatgca
19261	tegetggtgg	acgeetacat	caacaccggc	gacctacact	accgctccat	gctcctgggc
19321	gtcaatccct	tcaaccacca	ccgcaacgcg	gtacccaaa	agttctttgc	catcaagaac
19381	aacggccgct	acgtgccctt	ccacacccaa	gogocccaaa	tccacsagga	catcaacata
19441	ctgcttctgc	tccccggttc	Ctacacctac	gagtggaact	tccgcaagga	ccacttcaac
		~++~~~~~~~~	caaccacciu	Cucucuaca	gcgcccccg	
		~~~~~~~~~	caaccaccau	LCCLLCaacy	accace co	955
		+ ~~~~~~~~	ccancecatu	aucauucayy	cggccg	3
			CCTACCCLLC	Caucacaaca	uccoggggeee	
20281	. ggctccaccg	tagetecete	ctttatata	ataggggggg	tcaccgacct	gggtcagaac
20341	. cgcatcccat	tetecageaa	cccaegee	racatracct	ttgaggtgga	ccccatggat
20401	. atgctctatg	ccaactcggc	tatattaga	gacatgacca	togtcagagt	gcaccagccg
20461	. gagcccaccc	tcctctatct	celettegaa	gccccgacg	tggtcagagt	caacgctacc
20521	caccgcggcg	r tcatcgaggc	egtetacety	- cycacycccc	teteegeegg	gacctgggat
			TACACCLLCC	autacuayy	, cauge eges	- 2 - 2 - 2 - 2 - 2 -
2190	1 gtcaccttg	g ccttgctgg	g ctgctcctt	c aacgcgcgc	t gcccgttctc	gctggtcaca

FIG. 16A-6

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. .					antagnasas	attagastag
21961	tccatctcca	ccacgtggtc	cttgtggatc	accaccyccc	catycayata	cccgagccga
22021	ccctcgacat	cgcagcagcc	atgatcccac	agggcgcagc	cggtgcactc	ccagttetta
22081	tacacaatca	cactataact	gaagatgtaa	ccttgcaaca	ggcgacccat	gacggtgcta
221/1	aatgctttct	agataataaa	ggtcagttgc	agaccgcggg	cctcctcgtt	catccaggtc
22741	tggcacatct	tttaassast	ctcaatctac	traggratua	gettgtaage	atcococago
2220I	tggcacatet	tttggaagat	tteestesee	acattaataa	tatecatece	cttctcccag
22261	ccgctgtcga	cgcggtagcg	ttccatcage	acyttcatgg	cacccacgcc	actacacag
22321	gacgagacca	gaggcagact	cagggggttg	cgcacgttca	ggacaccggg	ggttgcaggt
22381	tegacgatge	gttttccgtc	cttgccttcc	ttcaacagaa	ccggaggctg	gctgaatccc
22441	acteceaega	ttacggcatc	ttcctggggc	atctcttcgt	cggggtctac	cttggtcaca
22501	tgcttggtct	ttctaactta	cttcttttt	ggagggctgt	ccacggggac	cacgtcctcc
22561	tcggaagacc	cadadcccac	cccctcatac	tttcaacact	tggtgggcag	aggaggtggt
22301	ggcggcgagg	agetestata	ctactccaac	gnatagcgcg	ccaacccata	accccaaaac
22621	ggcggcgagg	ggccccccc	cegereegge	agatagtgag	taccaccac	cattetttcc
22681	ggagtggcct	ctcgctccat	gaaccggcgc	acytectgat	egeegeegge	22020000
22741	taggggaaga	tggaggagca	gccgcgtaag	caggagcagg	aggaggactt	aaccacccac
22801	gagcaaccca	aaatcgagca	ggacctgggc	ttcgaagagc	cggctcgtct	agaaccccca
22861	caggatgaac	aggagcacga	gcaagacgca	ggccaggagg	agaccgacgc	tgggctccag
22921	catooctacc	tgggaggaga	ggaggatgtg	ctgctaaaac	acttgcagcg	ccaatccatc
22981	atcctccggg	acacctaac	cgaccggagc	gaaacccctc	tcagcgtcga	ggagctgtgt
22001	cgggcctacg	acctcaacct	cttctcgccg	cacatacccc	ccaaacqcca	gcccaacggc
23041	acctgcgagc	ageceaacc	teteaactte	tatecegtet	ttacaateee	cgaggccta
23TOT	acetgegage	ccaacccgcg	ccccaacccc	agatocco	tetectacea	caccasccac
23161	gccacctatc	acatetttt	caagaaccaa	aagatteetty		tateacttac
23221	acccgcgccg	acgcgctcct	cgctctgggg	cccggcgcgc	geatacetya	tategettee
23281	ctggaagagg	tgcccaagat	cttcgaaggg	ctcggtcggg	acgagacgcg	cgcggcaaac
23341	gctctgaaag	aaacagcaga	ggaagagggt	cacactagcg	ccctggtaga	gttggaaggc
23401	gacaacgcca	aactaaccat	gctcaagcgc	agcgtcgagc	tcacccactt	cgcctacccc
23461	gccgtcaacc	tecegeceaa	gatcatacat	cgcatcatgg	atcagctcat	catgccccac
22501	atcgaggccc	tccatcasac	tcannancan	caccccaaaa	acocccoocc	cataatcaac
23321	gacgagcagc	tegacgaaag	acteaggageag	cacascaca	aggetttgga	acagcggcgc
2358T	gacgagcagc	tegegegeteg	getegggaee	cycyaccccc	aggeeetgga	ccacttcttc
23641	aagctcatgc	rggccgrggr	cetggteace	ecegageteg	aatgcatgcg	cogcocccc
23701	agcgaccccg	agaccctgcg	taaggtcgag	gagaccctgc	actacacttt	caggcacggc
23761	ttcgtcaggc	aggcctgcaa	gatctccaac	gtggagctga	ccaacctggt	ctcatgcctg
23821	gggatcctgc	acgagaaccg	cctgggacag	accgtgctcc	actctactct	gaagggcgag
23881	acacatcaaa	actatgtccg	cgactgtgta	tttctcttta	tctgccacac	ctggcaagca
23941	gccatgggcg	totogcagca	atatetegag	gacgaaaatc	tgaaggagct	ggacaagctt
2//01	cttgctagaa	accttasass	actatagaca	ggcttcgacg	agcgcaccgt	cgcctcggac
24001	ctggccgaga	teatttta	adaecucctu	addcadacdc	tgaaaggggg	actaccaac
24001	ttcatgagcc	cogcccccc	agaacgcccg	cccectttce	ttctcgagcg	atctoggato
24121	ttcatgagcc	agagcatgtt	ycaaaactac	cycactetea	cccccgagcg	ccacasatat
24181	ctacccgcca	cctgcaacgc	attcccctcc	gaetttgtee	egergagera	ccgcgagtgt
24241	ccccgccgc	tgtggagcca	ctgctatctc	ttgcagctgg	ccaactacat	egectaceae
24301	tcggacgtga	tcgaggacgt	gagcggcgag	gggcttctcg	agtgccactg	ccgctgcaac
24361	ctatactccc	cacaccactc	cctggtctgc	aacccccagc	ttctgagcga	gacccaggtc
24421	atcootacct	tcgagctgca	aggtccgcag	gagtccaccg	ctccgctgaa	actcacgccg
24481	gggttgtgga	cttccgcgta	cctgcgcaaa	tttgtacccg	aggactacca	cgcccatgaa
24541	ataaagttct	traaggarca	atcococcca	cagcacgcgg	atctcacqqc	ctgcgtcatc
24541	acccagggcg	ccatcatcac	ccaattgcac	accatccaaa	aatcccccca	agagtttctt
24001	ctaaaaaagg	atacacacac	ctacctggac	cccsascaa	acasaatact	caacccgggt
24001	Ctaaaaaagg	grayayyyyr	ccaccoggac	ccccagacgg	gegaggegee	agreement
24721	ctccccagc	atgccgagga	agaagcagga	geegetagtg	gagcagacgg	aagaagaatg
24781	ggacagccag	gcagaggagg	acgaatggga	ggaggagaca	gaggaggaag	aattggaaga
24841	ggtggaagag	gagcaggaaa	cagagcagcc	cgtcgccgca	ccatccgcgc	cggcagcccc
24901	gccggtcacg	gatacaacct	ccacagetee	ggccaagcct	cctcgtagat	gggatcgagt
24961	gaagggtgac	ggtaagcacg	agcggcaggg	ctaccgatca	tggagggtcc	acaaagccgc
25021	gatcatcgcc	tocttocaag	actocooggo	gaacatcgct	ttcgcccgcc	gctacctgct
25021	cttccaccac	addatasacs	tecececaa	catattacat	tactaccotc	accttcacag
22001	ataacaacaa	222202000	anantencea	daddaddcct	gaggatege	gcgaacgagc
72T#T	ccaayaaaaa	gcaagcaaga	22222222	tottoooss	tetttatee	atttttaaa
25201	cctcgaccac	cagggagctg	ayyaaccyya	to the court of th		atttttcagc
25261	agagtcgagg	tcagcagcaa	gaactgaaag	taaaaaaccg	gcccccgcgc	tegeteacce
25321	gcagttgctt	gtaccacaaa	aacgaagatc	agctgcagcg	cactctcgaa	gacgccgagg
25381	ctctgttcca	caagtactgc	gcgctcactc	ttaaagacta	aggcgcgccc	acccggaaaa
25441	aaggcgggaa	ttacctcatc	gccaccatga	gcaaggagat	tcccacccct	tacatgtgga
25501	octatcaocc	ccagatgggc	ctggccgcgg	gcgcctccca	ggactactcc	acccgcatga
25561	actooctcan	taccaacac	togatgatct	cacgggtcaa	cggggtccat	aaccatcgaa
		-559	-5 -5			• •

FIG. 16A-7

25621	accagatatt	gttggagcag	gcggcggtca	cctccacgcc	cagggcaaag	ctcaacccgc
.25681	gtaattggcc	ctccaccctq	gtgtatcagg	aaatccccgg	gccgactacc	gtactacttc
25741	cacataacac	actooccoaa	atccacataa	ctaactcagg	tgtccagctg	gccggcggcg
25801	cttcccggtg	cccactccac	ccacaatcgg	gtataaaaac	cctggtgatc	cgaggcagag
25861	gcacacagct	caacgacgag	ttaataaact	cttcgatcgg	tctgcgaccg	gacggagtgt
25001	tccaactagc	cadeaccada	agatcgtcct	tcactcccaa	ccaggcctac	ctgaccttgc
23321	agagcagctc	ttcccacct	cactccaaa	gcatcggaac	cctccagttc	gtggaggagt
7330T	ttgtgccctc	catchacttc	sacccettet	cadatcacc	aggeetetae	ccggacgagt
20041	ttataccgaa	gttccacccc	atracacaan	caataascaa	ctacgactga	atgtcccatg
70T0T	gtgactcggc	tanagatagat	canttanda	atctggacca	ctaccaccac	ctacactact
7010T	tcgcccggga	cageteegee	ctcatctact	ttgagtttcc	cdaddadcac	cccaacggcc
70777	ctgcacacgg	gagetgegga	accetace	acaccacca	gtctcacctg	gtcaggttct
7078T	tcacccagca	agreetteete	accgragagg	accadaga	taccacctac	accotctact
26341	gcatctgtcc	accetteetg	ttaataata	accyggage	tactctttgt	ggtgagttta
26401	ataaaagctg	taccccgaag	attattaga	attection	atcatcasat	caacaagagg
26461	atcaacttca	aactaagaac	cettettgga	tttacctgca	accaccacaa	gaagtacatc
26521	atctagtttt	cctttgagga	toststagg	atagggagg	cctcctccaa	caacaatatt
2658T	ctcctaccta	atcacaacac	cactctagta	acetteteac	ttaaaaaaaa	aaagctaatt
26641	cttcatcgcc	acaateteac	caguggacia	accttttttag	acadaagggc	cttccacagt
26701	cttcatcgcc	ctattgtaga	aggaacttac	cagugucaga	cacasacate	taaccttctt
26761	ttcactttgg	tgaacgttac	eggeageage	acayetyete	ttccatctct	aacagagggt
26821	tctgatacta	acaaacctcg	tgtcggaggt	gagetttggg	tagtasttag	tacagaggge
26881	gggagttcta	ttgaagtggt	tgggtatttg	accutagggg	totttatete	ctgggtgcata
26941	gcagtgctgt	atcaacttcc	ttgctgggtc	gaaatcaggg	cattlatety	ctgggtcaga
27001	cattgtgggg	aggaaccatg	aaggggctct	tgctgattat	ecetteeetg	graggaggra
27061	tgctgtcatg	ccacgaacag	ccacgatgta	acattaccac	aggcaargag	aggaacgact
27121	gctctgtagt	tatcaaatgc	gagcaccatt	gtcctctcaa	catcacattc	aayaacaaga
27181	ccatgggaaa	tgtatgggtg	ggattctggc	aaccaggaga	tgagcagaac	cacacygica
27241	ctgtccatgg	tagcgatggc	aatcacactt	teggttteaa	attcattitt	gaagttatgt
27301	gtgatatcac	actacatgtg	gctagacttc	arggerrgrg	geceetace	aaggagaaca
27361	tggtgggttt	ttctttggct	tttgtgatca	tggcctgctt	gatgtcaggt	ctgctggtag
27421	gggctctagt	gtggtttctg	aaacgcaagc	ccaggtacgg	aaatgaggag	aaggaaaaat
27481	tgctataaat	tetttttctc	ttcgcacaac	catgaataca	gtgttccgta	tegtgetget
27541	ctctcttctt	gtagctttcg	gtcaggcagg	aattcatatt	attaatgcta	catggtggga
27601	taatataact	ttagtgggac	cctcagatac	tccagttacc	tggtatgatg	gcaagggatt
27661	gcaattttgt	gacggaagta	cagttaagaa	tccgcagatc	agacatactt	gtaatgatca
27721	aaacttaact	ctgattcatg	ttaacaaaac	ccatgaaaga	acatacatgg	gttacagaca
27781	tgacagtaag	ggaaaagtag	actataaggt	tacagtcatt	ccacctcctc	ctgctactgt
27841	aaagccacaa	ccagatccag	aaaatgtctt	tgtttatatg	ggaaataatg	taactttagt
27901	tagaceteca	ggaattccag	ttagttggta	ttatcataat	ggcacacagt	tctgcgatgg
27961	agataaaatt	attcatccag	aattcaacca	cacctgtgat	aaacaaaacc	ttacactgct
28021	gtttgtaaac	tttacacatg	atggaggcta	tcttggattc	aattacaaag	gtactcagag
28081	aattcagtat	gaggttatag	ttttagatcg	atttccaaat	tctggtcaga	tgaaaattga
28141	agaacaaagt	gaggaaacag	aacagaaaca	tactgagcat	aataaggctg	gacaaaagca
28201	gggtatagat	acaaatcaaa	agaaagctaa	taacagacaa	aagccatctc	aaaggccatc
28261	aagaagacgg	ccgacaaaca	ctcctgagac	aaaacaactt	acagtgtcta	ttgggtctaa
28321	cttaacttta	attaatccaa	atggaaaagt	cacttggtat	gatggtgatt	taaaaagacc
28381	atgtgaagaa	caaaactata	ggcttccaca	tcagtgtagt	gctcagaact	taactttaat
28441	taatotaact	aaatctcatg	agggaactta	ctatggcact	aatgacaaag	acgaaagcaa
28501	aagatacaga	gtgaaagtga	acactacaaa	ttctcaagct	gtaaaaatta	acccatatac
28561	cagacctact	actcctgatc	agaaacacag	atttgaatta	caaattgaaa	ataatgcaaa
28621	tgatgaagaa	tcaaaaattc	catctactac	tgtggcaatc	gtggtgggag	tgattgcggg
28681	cttcataact	ataatcatto	tcattctgtg	ctacatctgc	tgccgcaagc	gtcccagggc
28741	atacaatcat	atggtagacc	cactactcag	cttctcttac	tgagactcag	tcactttcat
28801	ttcagaacca	tgaaggettt	cacagettge	gttctgttta	. acataatcac	acttagtgta
28861	gctgcaaatg	gttttaaaca	tgttaatgtt	accagattaa	. gtaatgtaac	actgacagga
28921	gctggaatta	atactacato	gacagggtat	tttaatgagg	gtccaaaagg	aaaaaatggg
28981	tagatgaata	tttgcacatg	gggcgatcct	agatatgtgt	gccatggaaa	. tagcagtact
29041	attactaatc	ttacagttgt	ggcacttcta	aatttaacca	ctaacagaag	atttaaagca
29101	gaaagtttta	ctagtaacga	tggttatgaa	actaccagtg	caaaatttta	. tgaaattaaa
29161	attattgagg	ttccaacaac	tagagcaccc	accacagtta	ggacaacaca	. gcctaccact
29221	gtgcccacta	cacatccaac	caccacagte	agtacaacta	ttgagaccac	tactcatact
1 J 12 11 1	5 - 5 - Coucia				- -	

FIG. 16A-8

		acacaacagt	aceastact	actitattoa	ttagattttt	actgagagga
29281	acacagetag	acacaacagt	geagaacaee	acctcaagtg	ccttcagcag	cactgcaaat
29341	aatgaaagta	ctactgaaca	gacagaggct		tratraatra	acageettae
29401	ttaacttcgc	ttgcttggac	taatgaaacc	ggagtattata	gatgatte	tettacaatt
			PACTETECTO	millula La	uuacccccc	ccccgcggcc
20521	abbababaab	++~+~+~~~	caaaoccada	uauaaattta	ggcggcccac	a cacaggee
20641	tttagagtat	aataatcaac	catgattcct	aggttettee	tatttaatat	cccgccccgc
00701		+ atatactac	crrcacaacc	ullilacaca	CCCCGCCGG	0090000
00761		catacctcct	ctttacccta	ctaacctqca	ectgegreeg	caycaccycc
00001		teacetttet	dcadctcatc	gactggtgt	gegegeeta	Caaccaccac
00001		cccaatacac	ggacgagaac	gtagccagaa	ttttaayytt	cattegatea
00041		actcatacta	atatecetee	tatcccctgc	Colligioadu	tttgctgatt
20001		assistanca.	gacatatoga.	atttcttaga	Liguration	yayaaaaccy
20061		ctattacttc	ataattatta.	gggtagtcat	ggtetgetta	Lycaccece
3000T	atatgeette	gatctacccc	tottttaatc	ttggctggaa	ctctgttgag	gcattcacat
30121	ttgccattat	aaacagttca	ctacctcca	caccaccacc	cacaccqcct	ccccgcagaa
30181	acacactaga	tatgattcag	tagttagag	agececetee	ccaacccct	tccactgtta
30241	atcagttccc	cataaccggc		gececetaa	acctcgagat	ggacggccag
30301	gctactttca	cataaccggc	ggegatgact	gaccaccegg	accedagaca	ggccgccaag
30361	gcctccgagc	agcgcatcct	geaactgege	tagaagaga	ageaggageg	cctggtcaag
30421	gagctcctcg	atgccatcaa	catccaccag	tycaayaayy	aggategeet	cacctataaa
30481	caggcaaaga	tcacctacga	gctcgtgtcc	ggeggeaage	toppogat	agtcatcacc
20541	attacaccacc	SSSSSSSS	atteacetae	atggtgggcg	Leaaceccat	agicalcace
20001		acasascess.	caactacatc	cactoctcci	gcgaaagccc	cgagtgtatt
20661	+-atacatac	traamarret	ttacagactc	cacaacctcc	tececatgaa	Cigalguiga
20721	++	aaaaaccaat	caaacccttc	cccaattact	Cataayaata	aaccaccgga
20201		caataaadat	cacttactto	aaatctgaaa	gtatgtetet	ggtgtagttg
20041		actemmasee	ctcctcccag	ctctddtact	ccaqtccccg	gcgggcggcg
20001		acaccttcaa	accoratotca	aattcctddt	CCacaalll	Callycele
20061		tannenees	ccaaataaaa	gatgacttca	accccgtcta	CCCCtatggc
21021	+ a concorona	atcadaatat	CCCCTTCCTT	actedecet	Lugitude	cgacggaccc
24004		ascetacaat.	cctatcactc	aaactddctd	accedatege	Caccaccacc
31081	Cadadctttt	cactcaaggt	Tagagagaat	cttactatta	aaaaagatag	tggaaatcta
31141	ggggatgttt	ctaaggctcc	cttocaactt	acaactgata	aacagttgga	aattgcactg
31201	aaggtgaacc	ttgaagtcag	taatoocaao	cttoocataa	aagcaggtca	togattgaaa
31261	gcttatccat	aaattgctgg	thereseas	ttaacaaata	cacttataat	tttgactgga
31321	gtcattgaca	aaattgctgg	tetygaagge	actuatuuut	caadtagagg	agttggtata
31381	aaaggaatag	gtactgaaaa	tettyaaaac	tattttatt	aaaaaaataa	tttagttgct
31441	aacgtaagac	ttgctaaaga	tggaggtetg	telligata	addagggtgd	cccaaattat
31501	tggaataaac	atgatgacag	acgcactcta	tggacaactc	topacceate	taggagtess
31561	acaatcgatc	aggaaaggga	ttcaaagctc	accttagtat	taacaaaacy	roscastaat
21621		atototott	acttottota	aaaggaaaat	Llaylaacal	aaacaacaac
21601		ctcataaaaa	aatcacagta	aagctacttt	ttaatgaaaa	gygagtatta
21711	atamamantt	-coacacttaa	gaaagaatat	tggaactaca	gaaatgataa	Licializata
21001	tatasaccet	atoataatoc	agttccttt	atgccaaaca	taaaagctta	LUCLAAACCE
21061	2442424242	cttcaactaa	accadaadat	aaaaaaagty	Cigilaaaay	acacactycy
24021		- atattanaaa	cttoccagat	aaaactuttu	ttataactat	Laagiilaai
21001		, aatataatta	ttcaattacc	ELLGAALICA	Calyyycaaa	aaccucuguu
22041		· ++~=++~~+~	ctctttacc	LECTICCIALA	Ligiticaaya	aaatgaggac
20101			aaaatgaatt	catotatett	. Callyalll	, Lacactagea
20161		~+c+cccacc	accadeceat	ttcacagtqt	aaacgattct	, cicagcacyy
20221		+-~~~~==t	attetaatta	atacaaaaa	tugacttygg	gittataatt
20001			caaacddddd	tcaataatta	agatgaagcc	giccicigaa
32281	L cacacague	. cctggcgagc	ecactersace	atcacagtet	ggtgaaacga	gaagaacgca
32341	aagtcatcu	ayegggeee	acagtocats	tatacctctc	catcagege	ctcaacagtc
32401	cagattcata	ctcggaaaac	. aggatgggtt	. egagaaaaa	gagatracaa	gtctctctga
32461	L tctgccgccg	i addcrcaara	- eggetgetge	. tootootoo	, gggaccacac	r caccocatco
3252	L ctatgatcco	cacageette	agcatcagto	. coccagatact	. caccatatta	caccgcatcc
2050		- astattetes	cagtaagtgg	. accacataai	, caccalgile	Liciagoayou
2064		· ~~t~ctccan	r ccaaaaactca	Lattaaaaa	. gatqqaaccc	; acytyactat
20701	1+	- accacactat	· atcadatdcc	toccccicat	: qaacacact	Cucatataca
2076	1	- aaacstatct	· ctottcacaa	l tetgaeggta	i ccagggaaag	, cyclygliga
2202	1	, atasataact	· ctcctdaaco	: acacqqcca	i cadddidcci	, cecycecyae
3288	1 actgcaggg	a gcccggggat	gaacagtggc	: aatgcaggat	ccagcgctcg	g tacccgctca

FIG. 16A-9

32941	ccatctgagc	tctcaccaag	tccagggtag	cggggcacag	gcacactgac	atacatettt
33001	ttaaaatttt	tatttcctct	ggagtcaaga	tcatatccca	ggggactgga	aactcttgga
33061	ocaoootaaa	accaacaaca	catggtaatc	cacggacaga	acttacatta	tgataatctg
33121	catgatcaca	atcaggcaac	aggggatgtt	gttcagtcag	tgaagccctg	gtttcctcat
33181	cagategtag	taaacgggcc	ctgcgatatg	gatgatggcg	gagcgagctg	gattgaatct
33241	cogtttocat	tgtagtggat	tctcttgcgt	accttgtcgt	acttctgcca	gcagaaatgg
33301	gcccttgaac	agcagatacc	cctcctgcgg	ccgtcctttc	gctgctgccg	ctcagtcatc
33361	caactgaagt	acatccattc	tcgaagattc	tggagaagtt	cctctgcatc	tgatgaaaca
33421	aaaaacccqt	ccatgcgaat	tcccctcatc	acatcagcca	ggactctgta	ggccatcccc
33481	atccaqttaa	tgctgccttg	tctatcattc	agagggggcg	gtggcaggat	tggaagaacc
33541	atttttattc	caaacggtct	cgaaggacga	taaagtgcaa	gtcacgcagg	tgacagcgtt
33601	cccctccqct	gtgctggtgg	aaacagacag	ccaggtcaaa	acccactcta	ttttcaaggt
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33901	tagaaacaga	tectactact	ccaccacctg	cagcgtgttc	aaaacaacaa	gattcaataa
33961	gattctgccc	teegeeetga	gctcgcgcct	caatgtcagc	tgcaaaaaat	cacttaagtc
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34081	ggaaaacttt	aatgctccaa	agctagcacc	caaaaactgc	atgctggaat	aagctctctt
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34201	catttqcqta	atagaaaagt	cctgtaaata	agtcactagg	accccaggga	ccacaatgtg
34261	gtagcttaca	ccgcgtcgct	gaagcatggt	tagtagagat	gagagtctga	aaaacagaaa
34321	gcatgcacta	aactaaggtg	gctattttca	ctgaaggaaa	aatcactctc	tccaacaaca
34381	gggtacccac	tgggtggccc	ttgcggacat	acaaaaatcg	gtccgtgtga	ttaaaaagca
34441	gcacagtaag	ttcctgtctt	cttccggcaa	aaatcacatc	ggactgggtt	agtatgtccc
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34621	aaaaattata	tctattgcta	gtcccttcct	ggacgggagc	aatccctcca	ggactatcta
34681	tgaaagcata	cagagattca	gccatagctc	agcccgctta	ccagtagaca	gagagcacag
34741	cagtacaage	gccaacagca	gcgactgact	acccactgac	ccagctccct	atttaaaggc
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35041	tagccgtgcg	tcgtgacgtc	atttgcatca	tcttctctcg	tccaatcagc	gctggccccg
35101	ccctaaattc	aaaagctcat	ttgcatgtta	acttttgttt	actttgtggg	gtatattatt
	gatgatc					
CEO TI	กัฬก• ร					

Grp	Vaccine	Monkey	P	re	W	k 4	W	K 8	Wk	12
•	at Wk 0, Wk 4	, ID	Mock	Gag	Mock	Gag	Mock	Gag	Mock	Gag
1	Ad24AE 1gggAOrf6Ad5Orf6	00C072	3	4	4	381	3	150	3	68
•	10^11 vp	00C178	3	3	1	559	ו	743	0	635
		00C222	0	3	1 1	369	1	753	0	670
		00D011	1	9	9	211	4	273	0	520
•		00D023	0	6	0	295	1	459	1	368
		00D031	15	5	10 .	103	1	101	1	40
2	Ad24AE 1gogAOrl6Ad5Orl6	99C168	4	6	0	118	5	241	3	209
-	10^10 vp	99C170	10	5	5	241	3	141	9	103
	10-10-49	99C173	1	3	ŏ	23	ŏ	14	Ō	21
3	Ad24AE 1gcgAE4Ad5Orf6	99C154	0	3	0	93	0	60	1	53
•	10^10 vp	99C158	1	0	1	141	0	101	1	120
		99C177	0	0	0	45	0	39	0	79
4	MRKAd5-HIVgag	00C018	1	5	13	1025	0	824	3	753
	10^11 vp	00C034	0	4	5	219	5	404	0	491
	•	00C058	4	4	3	1086	0	440	0	439
5	MRKAd5-HIVgag	99C218	0	3	5	2500	0	1580	10	1655
•	10^10 vp	99C227	6	1	4	529	5	365	5	1004
		99D185	ND	ND	0	425	0	310	0	271

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Vaccine	Monkey	Gag-Specific (Wk 12)		
at Wk 0, Wk 4	ID	%CD4	%CD8	
Ad24∆E 1gag∆Orf6Ad5Orf6	00C072	0.02	0.02	
10^11 vp	00C178	0.05	0.38	
	00C222	0.02	0.40	
	00D011	0.02	0.27	
	00D023	0.01	0.11	
	00D031	0.01	0.01	
MRKAd5-HIVgag	00C018	0.05	0.41	
10^11 vp	00C034	0.06	0.18	
_	00C058	0.02	0.28	

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Grp	Vaccine at Wk 0, Wk 4	Monkey ID	Wk 4	WK 8
1	Ad24AE 1 gogAOrf6Ad5Orf6	00C072	<10	77
	10^11 vp	00C178	<10	26
	•	00C222	<10	423
		00D011	<10	98
		00D023	<10	<10
		00D031	<10	<10
2	Ad24AE Lacas Orfe Ad5Orfe	99C168	-10	<10
2	Ad24∆E1gcg∆Orf6Ad5Orf6	99C170	<10	
	10^10 vp	99C170	<10	<10
		990173	<10	<10
3	Ad24ΔE 1gagΔE4Ad5Orf6	99C154	<10	<10
	10^10 vp	99C158	<10	<10
		99C177	<10	<10
4	MRKAd5-HIVgag	00C018	34	1017
	10^11 vp	00C034	14	423
	•	00C058	46	934
	MDKA dE LIVana	000018	- 00	
5	MRKAd5-HIVgag	99C218	20	99 767
	10^10 vp	99C227 99D185	40 17	767
	•	כסו טפצ	17	342

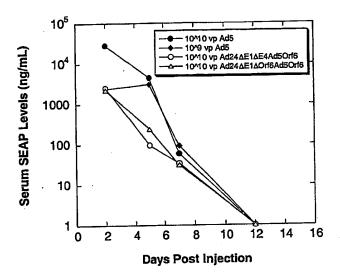


FIG. 20

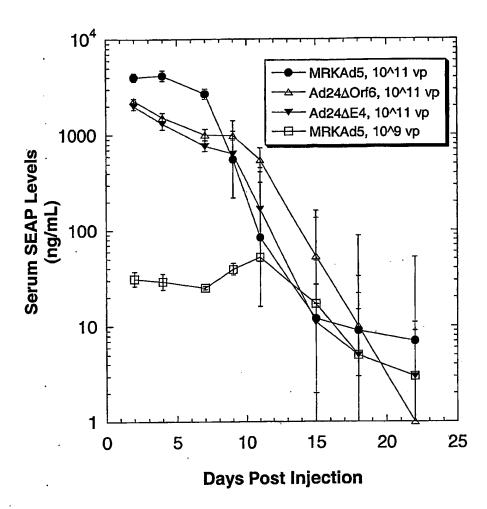
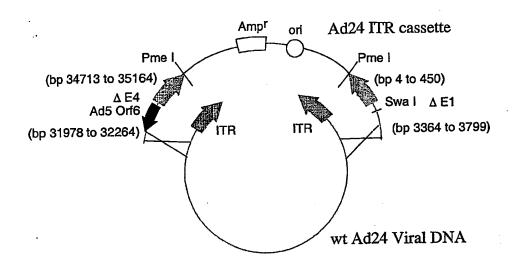


FIG. 21

40/59.

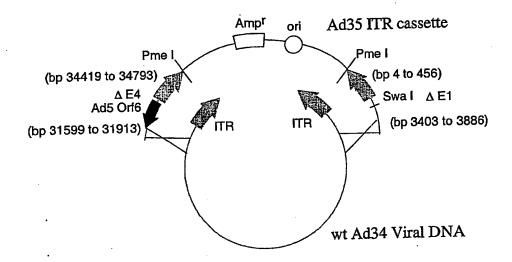


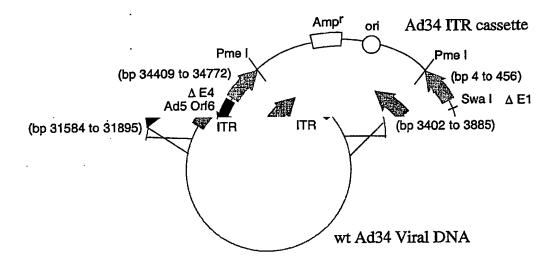
Animal	Prime (Wk 0, 4, 26)) Boost (Wk 56)		Pre		Primeb		Pre-Boost ^e		Post-Boost ^d	
	V (Mock*	Gag*	Mock	Gag	Mock	Gag	Mock	Gag	
Monkey 1	10° vp MRKAd5-gag	1011 vp Ad24ΔE1gagΔOrf6Ad5Orf6	18	16	1	244	3	74	3	1235	
Monkey 2	107 vp MRKAd5-gag	10 ¹¹ vp Ad24ΔE1gagΔOrf8Ad5Orf6	10	9	4	83	0	18	0	856	
Monkey 3	10° vp MRKAd6-gag	1011 vp Ad24ΔE1gagAOrf8Ad5Orf6	1	1	0	219	9	69	0	703	
Monkey 4	10 ⁷ vp MRKAd8-gag	1011 vp Ad244E1gag4Orf6Ad5Orf6	1	1	3	59	1	20	0.	419	
Mankey 5	none	1011 vp Ad24ΔE1gagΔOrf6Ad5Orf8	3	4	ND°	ND	ND	ND	4	558	
Monkey 6	none	1011 vp Ad24AE1gag&Orf6Ad5Orf6	0	3	ND	ND	ND	ND	1	295	
Monkey 7	none	1011 vp Ad24AE1gagAOrf8Ad5Orf8	1	9	ND	ND	ND	ND	9	103 ~	
Monkey 8	none	1011 vp Ad244E1gag&Orf8Ad5Orf6	3	3	ND	ND	ND	ND	1	381	
Monkey 9	none	1011 vp Ad24AE1gagAOrf8Ad5Orf6	0	6	ΝD	ND	ND	ND	0	369	
Monkey 10	лопе	1011 vp Ad24AE1gagAOrI6Ad5OrI6	15	5	ND	ND	ND	ND	10	211	

		Boost (Wk 56)	Gag-Specific T cells (Wk 60)		
Animal	Prime (Wk 0, 4, 26)	DOOS! (HER 30)	%CD4	%CD8	
		1011 vp Ad24ΔE1gagΔOrf6Ad5Orf6	0.06	0.37	
Monkey 1	109 vp MRKAd5-gag	10 ¹¹ vp Ad24ΔE1gagΔOrf6Ad5Orf6	0.01	0.56	
Monkey 2	107 vp MRKAd5-gag	10" Vp Ad24AE1gayAOrtoAd5Orto	0.07	0.06	
Monkey 3	109 vp MRKAd6-gag	10 ¹¹ vp Ad24ΔE1gagΔOrf6Ad5Orf6 10 ¹¹ vp Ad24ΔE1gagΔOrf6Ad5Orf6	0.04	0.20	
Mankovid	10 ⁷ vp MRKAd6-gag	10" VP Ad24AE IgagAONOAGSONG	0.0-1		

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Animal	Prime (Wk 0, 4)	Boost (Wk 24)	Pre		Prime ^b		Pre-Boost*		Post-Boost ^d	
	, , , , , , , , , , , , , , , , , , ,	· · ·	Mock ⁿ	Gag"	Mock	Gag	Mock	Gag	Mock	Gag
Mankey 11	1011 vp Ad24∆E1gag∆Orf8Ad5Orf8	107 vp MRKAd5-gag	3	4	3	150	4	28	0	188
Monkey 12	1011 vp Ad24AE1gagAOrf8Ad5Orf6	107 vp MRKAd5-gag	1 0	3	1	753	4	554	0	1029
Monkey 13	1011 vp Ad244E1gagAOrl6Ad5Orl6	10 ⁷ vp MRKAd5-gag	1	9	4	273	0	370	0	1520
Monkey 14	none	10 ⁷ vp MRKAd5-gag	0	0	ND*	ND	ND	ND	4	94
Monkey 15	none	107 vp MRKAd5-gag	0	0	ND ·	ND	ND	ND	1	168
Monkey 16	none	107 vo MRKAd5-gag	8	3	ND	ND	ND_	ND	8	149





-	catcatcaat	astatacett	atagatggaa	tagtaccaat	atgtaaatga	ggtgatttta
181	acgcataaaa aaatgaggta	aggettttt	t-acygaa	assattacta	atttgcgcgc	gaaaactgaa
.241	aaatgaggta	gttttgaccg	gatgeaagtg	tttataacaa	agtagagtat	ttattcaggg
301	aaatgaggta tgaggaagtg	tttttctgaa	taatgtggta	tttacggcag	catattttt	acctgaattt
			++sameerna	maaacutuca	acaq cagaaa	~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~
			actraanoro	ACULCAALAL	CCG CG CGCGC	9-9
			taattaactt	ai addadctu	ucccccccc	
2101	aagaggacaa	cccyayaycc	tacttactaa	atctacotcc	actggacggg	ataggggcgt taagtttaat
2161	gtctcctgaa	ctycaatyyy	ataataataa	toctagatet	gagttggctt	taagtttaat
2221	taagagggag	agggcatcta	geggeacega	r catgaggtc	cagaaagagg	gaagggatga
2281	gagtcgcaga	cgtcctgaaa	ccactoggeg	gcacgaggta	aaaacatott	ggttggagcc
2341	agtttctgta	ttgcaggaga	ttoose	ttetaccesa	atagetttga	ggcctgataa
2401	tgaggatgat	tgggaggtgg	CCalladaa	cocceetact	tottacatat	ggcctgataa
2461	acagtataag	attactagac	ggattaatat	. ccggaacgcc	agatoctoca	ctggaaatgg tgatggatat
2521	. ggctgaggtg	gtaatagata	Ctcaagacaa	ggcagttate	astattaaat	tgatggatat ttaggggaga
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204	. ageaggegg	aaa	•	: -		

2004	aaaaaaaaat			associatet	attattastt	taaaatcaac
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2227	tggtctccat	ggggccatat	cctccttccc	taacaaacaa	actatecata	teccentaga
228T	tggtetecat	gagttegt	terestran	tachaacag	ttattaataa	accasatata
5641	ctgattttac	aggcctcttc	tccagtggag	tgceteggte	LLCLLCGLac	aggaactery
5701	accactctga	tacaaaggcg	cgcgtccagg	ccagcacaaa	ggaggctatg	tgggagggt
5761	agcgatcgtt	gtcaaccagg	gggtccacct	tttccaaagt	atgcaaacac	atgtcaccct
5021	cttcaacatc	carraatoto	attogettat	aggtgtattt	cacataacct	agatececa
7021	ctgggggggt	caggaacgcg	gagattatt	actetteete	actotettee	ggatcgctgt
288T	ctgggggggt	acaaaagggg	geggeeeee	geceeeee	accocccc	agatogooga
5941	ccaggaacgt	cagetgttgg	ggtaggtatt	cccccccgaa	ggcgggcacg	accectgeac
6001	tcaggttgtc	agtttctaag	aacgaggagg	atttgatatt	gacagtgccg	gergagarge
6061	ctttcatgag	gttttcgtcc	atttggtcag	aaaacacaat	ttttttattg	tcaagtttgg
6121	toocaaatoa	tccatacagg	gcgttggata	aaagtttggc	aatggatcgc	atggtttggt
6181	tcttttcctt	atecacacac	tetttggcag	cgatgttgag	ttggacatac	tegegtgeta
6241	ggcacttcca	ttcccccaaa	atanttotca	attcatctcc	cacgattete	acttoccacc
0241	ggcacttcca		acageegeea	testaccogg	ctogasters	aggagetect
630T	ctcgattatg	caaggtaatt	adattedatae	tggtggccac	cucycucya	aggggcccgc
6361	tggtccaaca	gagcctacct	cctttcctag	aacagaaagg	gggaagtggg	tctagcataa
6421	gttcatcggg	agggtctgca	tccatggtaa	agattcccgg	aagtaaatcc	ttatcaaaat
6481	agctgatggg	agtggggtca	tctaaggcca	tttgccattc	tcgagctgcc	agtgcacgct
65/1	catatgggtt	ageggggeen	ccccadaca	taggataggt	gagtgcagag	gcatacatgc
0341	catatyggtt	adggggactg	aterraters	casacataca	tatatacatt	gratagrate
POOT	cacagatgtc	acagacgcag	atgygateet	caaayacycc	-tatatagget	ggacagcacc
6661	gccccctct	gatacttgct	cgcacatagt	catatagttc	argrgargge	gclagcaacc
6721	ccggacccaa	gttggtgcga	ttgggttttt	ctgttctgta	gacaatctgg	cgaaagatgg
6781	cgtgagaatt	ggaagagatg	atagatettt	gaaaaatgtt	gaaatgggca	tgaggtagac
	ctacagagtc					
0041	tgacaagtac	-t-ct-acadag	coggeataas	atattatta	aataatataa	taacctaatt
990T	tgacaagtac	gcccagggcg	caytayttaa	gracticity	aacyacycca	thacetyget
6961	ggtttttctt	ttcccacagt	tcgcggttga	gaaggtattc	ttegegatee	ttccagtact
7021	cttctagcgg	aaacccgtct	ttgtctgcac	ggtaagatcc	tagcatgtag	aactgattaa
7081	ctgccttgta	agggcagcag	cccttctcta	cgggtagaga	gtatgcttga	gcagcttttc
71/1	gcagcgaagc	ataaataaaa	acasaatat	ctctgaccat	gactttgaga	aattootatt
1747	tgaagtccat	atoatasas-	gotocotott	coccacactta	gaagtotage	catttattat
/201	LyaagtCCat	gregteacag	golddolgit	cccagaguig	yaaytttatt	ambabasas
7261	aggcggggtt	gggcaaagcg	aaagtaacat	cgttgaagag	aatcttaccg	gerergggea
7321	taaaattgcg	agtgatgcgg	aaaggctgtg	gtacttccgc	tcgattgttg	atcacctggg
7381	cagctaggac	gatetegteg	aaaccottoa	tattatatac	tacgatgtat	aattctatga'
7//1	aacgcggcgt	accept the acc	tracretarch	tattgaggtg	atcasaggtt	aggtetatag
/441	aacycyycyt	goodettyacg	taresses	naticyayett	atanagett	agg couguag
7501	ggtcagataa	ggcgtagtgt	Legagageee	attegtgeag	grgaggarrt	ycarytagga
	atgatgacca					
7621	gctggccaat	tgccattttt	tctggagtga	cacagtagaa	ggttctgggg	tcttgttgcc
7681	atcgatccca	ctttagttta	atggctagat	catagaccat	gttgacgaga	cgctcttctc
77/1	ctgagagttt	catnaccanc	atgaaaggaa	ctagttgttt	gccaaaggar	cccatccagg

FIG. 28A-2

			10/05			
7801	totaaottto	cacatcgtag	gtcaggaaga	gtctttctgt	gcgaggatga	gagccgatcg
7061		ペッセナナベクナベベ	caccagrigg	addattadtt	u c cua cg cg ~	
7001		MARKARACHAR T	CATECGEGEE	Latactuata	Cagacagecag	
T 0 0 1	++~~~~	aaattatata	teataataa	actutatet	gccccccg	-coaracre-
0041	h-n-~+~~~	TOOMSOOCE .	aacaartata	ECECULACIO	LLCLacaccc	9009000099
0101		++~+~+++	araaraarca .	Luctuacuau	CCCCCGCGGG	
01/1		MANAGEMANA	COCCCCCAA	ddaccauauc	gcgcaggccg	gagoogooo
0001		SACCTACAGE	CECAGGEEAG	TaddLaddua	Lagaagacca	~~~~~~~~~
0001	*********	accatacaaa	addecadae	dotactidat	Liccacagge	cegeegemg
0044		aaattaaaaa	atteceatate.	CEEEGGGCGC	Caccaccyca	ccccgaaaa
8401	ttcttttgat	cggtggtggc	tetettgett	erregranger	actantanta	gcacgtcggc
8461	cacaccaaac	ggaagcggtt ggcaggttct	gttccggacc	totaggearg	cttacataca	ccaccacgcg
8521	gccgcgcacg	tcttgtatct	ggtattgtgt	artasaaact	accooccco	tgagcttgaa
0644		arttraaran	aarcaarrec	dotatcutta	acqueagett	900000
0001		tasacanant	torcordata	ddcdatcttt	gccatgaact	gcccgaccc
07.71	+++-a+~a	agateteege	dacccdctct.	CECGACGGEG	guugugge	Cacoggaga
0001		anttaggaga	argcagtcat	acccaccica	Lucagacyc	990090000
0001		teggaagtete	ttacacacat.	caccaccida	gugaggutaa	gccccacgog
0041	+-++~==~	accocataot	tacatagaca	ctdaaaaauu	Lagitgagig	cggcggcaac
0001		accascast	acatgatcca	ECOTECECAGE	ggcarrege	cgacaseges
00/1		aaddactcca	raaccecata	daadiccacu	gcaaaaccaa	uuuuu u j j j j
0101		ascscactca.	arrececce	dadaadacuu	atyayttty	
0101		aattaaaaa	crecedaar.	CECELCULUC	LULUULALUU	CCCCCCCCC
0041	+	+~++~~+	canacaaaaa	coaaaaaaaa	acacygogae	999
0201		caataataa	arcottcaat	dacctctccu	cggcggcggc	gcarggroom
0261		caaccattet	cacacaarca	cadadladaa	acaccyccyc	gcaccopco
0421		ctacacatt	crccarread	dadddadadd	gegetgatta	Cucacccaa
9481	taattggccc	gtagggactg	cgcgcagaga	cetgategeg	ratagacca	gtacggcttc
9541	aaacctttcg	acgaaagcgt	ctaaccagte	taggicacaa	ggtaggttga	catctcggga
9601	ttgtgggcgg	gggtggttat atgctgctgg	_grgrrcggrc	aaagtaggca	gttctaagac	ggcggatggt
9661	aggtgagacg	accaggtett	tagatecage	tractagata	cacaaacaat	tggccattcc
0701		tectracate	tagcaagatc	ELLULAGIAG	Lucigualya	gccgccccac
0041	asattat	tactcaccca	rrctgccatg	catacululu	aytttaaatt	cgcgcaccgg
0001	++-+-aa-a+	acceentrea	ctacgactct	ttcaacaaaa	atygutugut	gcaccaggga
00.01		tananatant	caaaatccac	aaaucuuluu	Laayyuuu	caccaacggc
10001	~+~~~~~	· ttaaccataa	craaccaatt	aactutctuu	Lyaccayyyc	gcacgagece
40001		34444444	ACCCCCCCCCC.	ulcaaauatu	Lagicy	49909090
40461		, <u> </u>	таааатосоо	caataattaa	cygtagagag	gccacegee
4 00 01		~~~~~~~~~~	CASAALCEEC	Caacalaauu	Lugicacage	Cg cagacg ca
10001		. <i>~~~~~~~~~~~</i>	- cracaacaa u	autauaaucu	Lydygadact	0909000909
40001		, ++~~~~+a~~~	acardaadta	. Ollicatiu ca	, 4400099000	94004905-5
4 0 2 0 1		, tasttastac	гстагадаса	Luuauaaaa	. uaaaycytte	49094005-
10441	ctccgtagcc	tggaggaacg	tgaacgggtt	gggtegeggt	actcccqtct	casccaacc
10501	tactcgagcc	ggccggagcc	geggetaacy	tygtattygt tttactaat	toccoaatgo	cagggaagtg
10561	tacaaaaatc	caggatacgg ttttttttg	aaccyayccy	gcatcccgto	r ctgcgacaga	tgcgtcccca
10621	agtcctattt	ccctcgcagc	accaccagac	acaaaaggct	gtccctgcaa	ctactgcaac
10981	acaacagccc	ageggtgegg	. agcagcaacc	ctatgatcto	gacttggaag	agggcgaagg
40001		. ~-~~~+~~~	CETCOCCCO	ucuucatee	Luductoduc	. cgaaaaaaaga
40001		- ~~~+>+~+^	CCCAACAGAC	CCLatttauc	ı uacayaayç	9090990500
10001		* acadettee	rotttaacuu	: uuuttuutu	Cuyuyuuuu	9
10001		· ttacaaaaaa	. addatttcda	i autuuatua	ı utgacayyy	t ccagcooss
44041		- ~+~~~+~~	, ccaaccttd:	. alcuucta	. uaacauacay	Lauraggaas
4 4 4 0 1		. ~~~~~~~~~	rraaraarca	LULUCHAAC	. Cccaccycc	, gugaagaag
		- ttastacatt	. cacaaaacii	. uatyyaayci	, allallugi	t accerace
44001		- ~~~~~~~~~	· rarrreraa:	. uulucaaca	, aucayayac	i acgaggeeee
44004		- atactesaes	rcaccdaac	: cuauuuuau	i tugutuguati	acceracea
44744		 artatratar 	t tacaaaaac	, dadeecquu	: Clydccyay	aggryggerge
44 404		- +aaatttta	acredodaaa	i dialiacyci	Liguagate	, acaagaceee
11111		~ atamamann	t acctdaacai	. adaluqqil	: Lacalycyc	ı iyalyılıyaa
112501		- atasacasta	t arcecoodd	. GLaccucaa	L uacayaary	, accycycygc
44704		_ ~~~~~~~~~~	t actraacco	i Cauduaacu	ı alulacayı	, tycaaayayc
11641	l tctaactgg	a gctggaaccg	g agggtgagaa	a LLactinga	c acygyayct	g acttgcagtg

11701	gcagcctagt	cgcagggctc	tgaacgccgc	gacggcagga	tgtgagcttc	cttacataga
11761	agaggcggat.	gaaggggagg	aggaagaggg	cgagtacttg	gaagactgat	ggcacaaccc
11821	atattttta	ctagatggaa	cagcaagcac	cggatcccgc	aatgcgggcg	gcgctgcaga
11881	accaaccatc	cggcattaac	tcctcggacg	attggaccca	ggccatgcaa	cgtatcatgg
11941	cattagacaac	tcgcaacccc	gaageettta	gacagcaacc	ccaggccaac	cgtctatcgg
12001	ccatcatca	agctgtagtg	cetteceast	ctaatcccac	tcatgagaag	atcctaacca
12001	toatoacoga	gttggtggag	aacaaaacta	ttcgtccaga	tgaggccgga	ctootataca
12001	ccgcgaacgc	agaacgcgtg	actcactaca	acadtadcaa	totocaaacc	aatttggacc
12121	atguitte	agatgtacgc	geeegeeatat	ctcagggggg	aaggttccag	cacaatacca
12101	graryaraac	gctggtggcg	ttasatactt	tettgagtag	tragectect	aatotococ
12241	accegggee	ggattatact	natttta	atactttaaa	actostoota	transantar
12301	gtggtcaaca	agtatatact	taggetagta	attacttctt	tracactace	anacangort
12361	ctcagagcga	agtatateag		accaccccc	agattata	agacagggcc
12421	tgcagacggt	aaatctgagc	caagetttta	aaaaccttaa	taggettg	caatattat
12481	ccccggtagg	agaaagagca	accgtgtcta	gettgttaae	rectantes	totttaccat
12541	tactgttggt	agctcctttc	accgacagcg	gragearega		caccigggic
12601	acctactaaa	cctgtatcgc	gaagccatag	ggcaaagtca	ggrggacgag	tagaectate
12661	aagaaattac	ccaagtcagt	cgcgctttgg	gacaggaaga	cactggcagt	ttggaageea
12721	ctctgaactt	cttgcttacc	aatcggtctc	aaaagatccc	tcctcaatat	gctcttactg
12781	cggaggagga	gaggatcctt	agatatgtgc	agcagagcgt	gggattgttt	ctgatgcaag
12841	agggggcaac	tccgactgca	gcactggaca	tgacagcgcg	aaatatggag	cccagcatgt
12901	atoccaotaa	ccgacctttc	attaacaaac	tgctggacta	cttgcacaga	gctgccgcta
12961	tgaactctga	ttatttcacc	aatgccatct	taaacccgca	ctggctgccc	ccacctggtt
13021	tctacacggg	cgaatatgac	atgcccgacc	ctaatgacgg	atttctgtgg	gacgacgtgg
13081	acagcgatgt	tttttcacct	ctttctgatc	atcgcacgtg	gaaaaaggaa	ggcggcgata
13141	gaatgcattc	ttctgcatcg	ctatccaaaa	tcattggtgc	taccgcggct	gagcccgagt
13201	ctacaeatca	ttttcctagt	ctaccetttt	ctctacacag	totacotage	agcgaagtgg
12261	atacaataaa	tcgcccgagt	ttaatggggg	aagaggagta	cctaaacgat	tecttectca
13201	grayaaraay	agaaaaaaat	ttaccasaca	atogagagaa	aagtttggtg	gataaaatga
13321	gaccggcaag	gacttatgct	coccaaaca	acggaacaga	tagactagaca	rrractacaa
13381	gtagatggaa	gacttatget	cayyattata	gagacgagcc	agglattata	tacascasta
13441	gtagagcgag	ccgtagacgc	cagegeeacg	acayacayay	gggtcttgtg	raggacgacg
13501	aggattcggc	cgatgatagc	agcgtattgg	acttgggtgg	gagaggaagg	ggcaacccgt
13561	ttgctcattt	gcgccctcgc	ttgggtggta	tgttgtaaaa	aaaaataaaa	aagaaaaac
13621	tcaccaaggc	catggcgacg	agcgtacgtt	catterrerr	tattatetgt	gtctagtata
13681	atgaggcgag	tcgtgctagg	cggagcggtg	gtgtatccgg	agggtcctcc	teettegtae
13741	gagagcgtga	tgcagcagca	gcaggcgacg	gcggtgatgc	aatccccact	ggaggctccc
13801	tttatacctc	cocoatacct	ggcacctacg	gagggcagaa	acagcattcg	ttactcggaa
13861	ctggcacctc	agtacgatac	caccaggttg	tatctggtgg	acaacaagtc	ggcggacatt
13921	acttctctaa	actatcagaa	tgaccacagc	aacttcttga	ccacggtggt	gcaaaacaat
13981	gactttaccc	ctacggaagc	cagcacccag	accattaact	ttgatgaacg	atcgcggtgg
14041	ggcggtcagc	taaaaaccat	catgcatact	aacatgccca	acgtgaacga	gtatatgttt
14101	agtaacaagt	tcaaagcgcg	tgtgatggtg	tccagaaaac	ctcctgaggg	tattagagta
14161	racrataatt		- 5 - 5 5 5 5			cg c cagag ca
14221		acoaccacaa	gcaagatatt	ctaaaatacg	agtggttcga	gtttactttg
1/281	ccadaaddca	actiticaca	gcaagatatt	ctaaaatacg	agtggttcga	gtttactttg
	ccagaaggca	acttttcggt	cactatgact	ctaaaatacg atcgacttga	agtggttcga tgaacaatgc	gtttactttg catcatagac
1/3/1	ccagaaggca aattacttga	acttttcggt aagtgggcag	cactatgact acagaatgga	ctaaaatacg atcgacttga gtgttggaaa	agtggttcga tgaacaatgc gtgacattgg	gtttactttg catcatagac tgttaagttc
14341	ccagaaggca aattacttga gacactagga	acttttcggt aagtgggcag acttcaagtt	cactatgact acagaatgga gggatgggat	ctaaaatacg atcgacttga gtgttggaaa ccagaaacta	agtggttcga tgaacaatgc gtgacattgg agttgatcat	gtttactttg catcatagac tgttaagttc gcctggggtt
14341 14401	ccagaaggca aattacttga gacactagga tacacctatg	acttttcggt aagtgggcag acttcaagtt aggccttcca	cactatgact acagaatgga gggatgggat tcctgacatc	ctaaaatacg atcgacttga gtgttggaaa ccagaaacta gtattgctgc	agtggttcga tgaacaatgc gtgacattgg agttgatcat ctggctgcgg	gtttactttg catcatagac tgttaagttc gcctggggtt agtggacttt
14341 14401 14461	ccagaaggca aattacttga gacactagga tacacctatg accgaaagcc	acttttcggt aagtgggcag acttcaagtt aggccttcca gtctgagcaa	cactatgact acagaatgga gggatgggat tcctgacatc ccttcttggc	ctaaaatacg atcgacttga gtgttggaaa ccagaaacta gtattgctgc attagaaaga	agtggttcga tgaacaatgc gtgacattgg agttgatcat ctggctgcgg aacacccatt	gtttactttg catcatagac tgttaagttc gcctggggtt agtggacttt ccaagagggt
14341 14401 14461 14521	ccagaaggca aattacttga gacactagga tacacctatg accgaaagcc	acttttcggt aagtgggcag acttcaagtt aggccttcca gtctgagcaa tgtatgagga	cactatgact acagaatgga gggatgggat tcctgacatc ccttcttggc tttagaagga	ctaaaatacg atcgacttga gtgttggaaa ccagaaacta gtattgctgc attagaaaga ggaaatattc	agtggttcga tgaacaatgc gtgacattgg agttgatcat ctggctgcgg aacacccatt cagccctttt	gtttactttg catcatagac tgttaagttc gcctggggtt agtggacttt ccaagagggt ggatgtagat
14341 14401 14461 14521 14581	ccagaaggca aattacttga gacactagga tacacctatg accgaaagcc tttaagatct gcttatgaga	acttttcggt aagtgggcag acttcaagtt aggccttcca gtctgagcaa tgtatgagga acagcaagaa	cactatgact acagaatgga gggatgggat tectgacatc ccttcttggc tttagaagga agatcaaaaa	ctaaaatacg atcgacttga gtgttggaaa ccagaaacta gtattgctgc attagaaga ggaaatattc gccaaaatag	agtggttcga tgaacaatgc gtgacattgg agttgatcat ctggctgcgg aacacccatt cagccctttt aagctgctgc	gtttactttg catcatagac tgttaagttc gcctggggtt agtggactt ccaagagggt ggatgtagat agaagctaaa
14341 14401 14461 14521 14581 14641	ccagaaggca aattacttga gacactagga tacacctatg accgaaagcc tttaagatct gcttatgaga gcaaacatag	acttttcggt aagtgggcag acttcaagtt aggccttcca gtctgagcaa tgtatgagga acagcaagaa ttgccaacga	cactatgact acagaatgga gggatgggat teetgacate cettettgge tttagaagga agatcaaaaa teeggtaagg	ctaaaatacg atcgacttga gtgttggaaa ccagaaacta gtattgctgc attagaaaga ggaaatattc gccaaaatag gtggctaacg	agtggttcga tgaacaatgc gtgacattgg agttgatcat ctggctgcgg aacacccatt cagccctttt aagctgctgc ctagtgaaat	gtttactttg catcatagac tgttaagttc gcctggggtt agtggacttt ccaagagggt ggatgtagat agaagctaaa caggggagac
14341 14401 14461 14521 14581 14641 14701	ccagaaggca aattacttga gacactagga tacacctatg accgaaagcc tttaagatct gcttatgaga gcaaacatag agttttgcca	acttttcggt aagtgggcag acttcaagtt aggccttcca gtctgagcaa tgtatgagga acagcaagaa ttgccaacga caacatccgt	cactatgact acagaatgga gggatgggat tectgacatc ccttcttggc tttagaagga agatcaaaaa tccggtaagg tccgactaag	ctaaaatacg atcgacttga gtgttggaaa ccagaaacta gtattgctgc attagaaaga ggaaatattc gccaaaatacg gtggctaacg gaatcattat	agtggttcga tgaacaatgc gtgacattgg agttgatcat ctggctgcgg aacacccatt cagcccttt aagctgctgc ctagtgaaat tggatgatgt	gtttactttg catcatagac tgttaagttc gcctggggtt agtggacttt ccaagagggt ggatgtagat agaagctaaa caggggagac gtctcaaaac
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14341 14401 14461 14521 14581 14641 14701 14761 14821	ccagaaggca aattacttga gacactagga tacacctatg accgaaagcc tttaagatct gcttatgaga gcaaacatag agttttgccga atagagtcaa gtgttggaag	acttttcggt aagtgggcag acttcaagtt aggcettcca gtctgagcaa tgtatgagga acagcaagaa ttgccaacga caacatccgt aactcactat ataaaatcaa	cactatgact acagaatgga gggatgggat tectgacate cettettgge tttagaagga agatcaaaaa tceggtaagg tcegactaaa taageetgtg caeggeetat	ctaaaatacg atcgacttga gtgttggaaa ccagaaacta gtattgctgc attagaaaga ggaaatattc gccaaaatacg gtggctaacg gaatcattat gaaaaagatg cgcagttggt	agtggttcga tgaacaatgc gtgacattgg agttgatcat ctggctgcgg aacacccatt cagccctttt aagctgctgc ctgatgaaat tggatgatgt gcaaaaacag acctttcgta	gtttactttg catcatagac tgttaagttc gcctggggtt agtggacttt ccaagagggt ggatgtagat agaagctagat agagggagaa gtctcaaaac aagttacaat caattatggc
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14341 14461 144521 14581 146701 14761 14881 14941 15001 15121 15181 15241 15301 15321	ccagaaggca aattacttga gacactagga tacacctatg accgaaagcc tttaagatct gcttatgaga gcaaacatag agttttgccg atagagttaa gtgttggaag gaccccgaaa gcgagcagg agcttctaca cacgtcttca accaccgtca agtatccggg gtgtacaagg ataaaaaaaaaa	acttttcggt aagtgggcag acttcaagtt aggccttcca gtctgagcaa atgtatgaga acagcaagaa ttgccaacga caacatccgt aactcactat ataaaatcaa aaggagtgcg tctagtaacta acgaacaagc acgctttcc gtgaaaacgt gagtccaacgt aactgggcat aatgtcggtt	cactatgact acagaatgga gggatgggat tcctgacatc ccttcttggc tttagaagga agatcaaaaa tccggtaagg tccgactaaa taagcctgtg cacggcctat ttcctggaca gcttccagac ccctgtggtg tgtgtactcc tgagaaccag tcctgctct tgtgaccgt agtcgcacg	ctaaaatacg atcgacttga gtgttggaaa ccagaaacta gtattgctga ggaaatattc gccaaaatag gtggctaacg gaatcattat gaaaaagatg cgcagttggt ttgctaaca atgatgcagg ggtgcagagc cagcagctcc attttaatcc acagatcacg actgacgcctt cagtaataac acgcttcttac	agtggttcga tgaacaatgc gtgacattgg agttgatcat ctggctgatcat cagcccttt aagccgttgc ctagtgaaat tggatgatgt gcaaaaacag accttcgta acctctagatgt atcctgtcac ttatgcccgt gccagtccac gtccgccgcac caagccgcac	gtttactttg catcatagac tgttaagttc gcctggggtt agtggacttt ccaagagggt ggatgtagat agaagctaaa caggggagac gtctcaaaac aagttacaat caactacggc ctctcccctttcaaag ctcgctcc cttttcaaag gcccacaatt gttgcgcagc ctgtccctac ttctacaaa gcctgcggc
14341 14461 14461 14521 14581 14641 14761 14761 14881 15001 15181 15181 15241 15361 15361 15481	ccagaaggca aattacttga gacactagga tacacctatg accgaaagcc tttaagaatct gcttatgaga gcaaacatag agttttgccg atagagttaa gtgttggaag gaccccgaaa gcggagcaag actagacaag agtttctaca acaccttca acaccgtct acaccgtca agtatccggg gtgtacaagg gatacaaga accagcagt	acttttcggt aagtgggcag acttcaagtt aggccttcca gtctgagcaa tgtatgagga acagcaagga caacatccgt aactcactat ataaaatcaa aaggagtgcg tctactggtc tcagtaacta acgaacaagc accgctttcc gtgaaaacgt gagtcggcat aattcggta	cactatgact acagaatgga gggatgggat tcctgacatc ccttcttggc tttagaagga agatcaaaaa tccggtaagg tccgactaaa taagcctgtg cacggctat ttcctggaca gcttccagac ccctgtggtg tgtgtactcc tgagaaccag tcctgctct tgtgaccgtt agtcgcaccg tattccagac	ctaaaatacg atcgacttga gtgttggaaa ccagaaacta gtattgctgc attagaaga ggaaatattc gccaaaatag gtggctaacg gaatcattat gaaaaagatg cgcagttggt ttgctcacca atgatgcagg ggtgcagagc cagcagctcc attttaatcc acagatcacg actgacgcca cgcgtcctt cagtaataac acgttctacc caagggccgc	agtggttcga tgaacaatgc gtgacattgg agttgatcat ctggctgcgg aacacccatt cagccctttt aagctgctgc ctagtgaaat tggatgatgt gcaaaaacag acctttcgta cctcagtgcac ttatgccgt gccagtccac gtccgcggc ggaccctgcc gacgccgcac caagccgcac accggttggg caacatcccg actcggt	gtttactttg catcatagac tgttaagttc gcctggggtt agtggactt ccaagagggt ggatgtagat agaagctaaa caggggagac gtctcaaaac aagttacaat caactactgcgc ctctttcaaag ctcgctacc gcccacaatt gttgcgcagc cttttctaaaa gtctgcgcg ttctgcgcgc ttctctacaa gtctgcgcg tgcgcaccact gttgcgcagc ctgtccctac ttctacaaa gtctgcgcg tgcgcgcgcg tgcgcgcgcgcgcgcgcg

FIG. 28A-4

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15601	atctactgtg	gacgcagtta cgaaggcgca	Ligadagigi	taaaaaaaact	accactocca	tacaaacaac
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12/01	~~~~~	gccgacatgg	cccaatcgcg	aagaggcaat	gtatactggg	tgcgtgacgc
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17401	atttttcate	ccaactgaac	-cgacacaggest	treattogag	cagtatetgg	agcgggctta
17461	teggcaegag	ccaactgaac	gggggcgccc	~~~~~~~~~	ttggaacagc	agtacaggac
17821	gcgtagatga	accgccitci	cacyayyaay	caacgaage	attacatcaa	cccatcacct
10001	. cgaaaceag	tctgaacagc	atcotoggto	taggcgtgca	. aagtgtaaaa	cgccgtcgct
18183	L cgccgtcaca	gcatcayayy	aaaaaaaggaa	taggoogis	treacaters	cogacaggat
18241	L aagatggcca	cccatcgat	getgetetaa	Lyggeacacc	. gggggggggg	cacctacttc
		+-?//////	aaarccracc	an Aucucus	, ,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	
1854	I ttctttgace	LLaggggege	. tassaaaaaa	ccaaatgcat	ctcagtggtt	ggataaggga
1860	l gcttacaaci	ccctggctcg ctggcctagt	Laaaggegee	. coadatget.	ataaaaaaa	agccaaaaaa
1000	1 totatocc	a acagaccca	a ctacattoo	ttcagagat	a acttcatcgg	g acttatgtac
1926	_ tataacagt	a ctggcaacat	- gagageact	tcttaccaa	c tettgetta	a ctctctgggc
1944	1 cgtgttatt	a gatactitag g aaaatcatg	g tgtggaaga	t gaadtteed	a accacage	

19501	ggtgtcggtc	cgcgaacaga	tagttacaag	gagattaagc	caaatggaga	ccaatctact
19561	tggacaaatg	tagacccaac	tggcagcagt	gaacttgcta	agggaaatcc	atttgccatg
19621	gaaattaacc	ttcaagccaa	tctatggcga	agtttccttt	attccaatgt	ggctctatat
19681	ctcccagact	cgtacaaata	caccccgtcc	aatgtcactc	ttccagaaaa	caaaaacacc
19741	tacgactaca	tgaacgggcg	ggtggtgccg	ccatctctag	tagacaccta	tgtgaacatt
19801	ggtgccaggt	ggtctctgga	toccatogac	aatqtcaacc	cattcaacca	ccaccgtaac
19861	actaacttac	gttaccgatc	catacttcta	ggtaacggac	attatatacc	tttccacata
19921	caagtgcctc	aaaaattctt	coctottaaa	aacctgctgc	ttctcccagg	ctcctacact
19981	tatgagtgga	actttaggaa	ggatgtaaac	atouttetac	agagttccct	cootaacoac
20041	ctacqqqqqq	atggcgccag	catcagtttt	acqaqcatca	acctctatoc	tacttttttc
20111	ccataacta	acaacaccgc	ttccaccctt	gaagccatgc	tacagaatga	caccaatgat
20161	cecacggeee	acgactacct	atctccacct	aacatoctct	accccattcc	taccaataca
		ccatttccat				
		aaaccaaaga				
		ctattcccta				
20401	gtttccatca	tgtttgactc	ttcagtgagc	tggcctggaa	atgacaggtt	actateteet
		aaataaagcg				
20521	atgaccaaag	actggttctt	ggtacagatg	ctcgccaact	acaacatcgg	ctatcagggc
		cagaaggata				
20641	atgagcaggc	aggtggttga	tgaggtcaat	tacaaagact	tcaaggccgt	cgccataccc
20701	taccaacaca	acaactctgg	ctttgtgggt	tacatggctc	cgaccatgcg	tcaaggtcaa
20761	ccctatcccg	ctaactatcc	ctatccactc	attggaacaa	ctgccgtaaa	tagtgttacg
20821	cagaaaaagt	tcttgtgtga	cagaaccatg	tggcgcatac	cgttctcaag	caacttcatg
20881	tctatgggag	cccttacaga	cttgggacag	aacatgctct	atgccaactc	agctcatgct
20941	ctggacatga	cctttgaggt	ggatcccatg	gatgagccca	ccctqcttta	tcttctcttc
21001	gaagttttcg	acgtggtcag	agtgcatcag	ccacaccaca	gcatcatcga	ggcagtctac
21061	ctacatacac	cgttctcggc	contaacoct	accacgtaag	aagettettg	cttcttgcaa
21121	acaccacto	caaccatggc	ctacaaatac	caaaacaact	ccadcdadca	agageteaga
21101	acagcagccg	aagacctggg	ttgcggaccc	tatttttaa	reaccetters	taancocttc
21241	gecattytee	tggcccccga	taagatagaa	tatacastta	tasatacccc	caagegeeee
21241	ceggggttea	rggeeeeega	caagetegee	teresseess	attatasasa	ctactacatt
21301	acggggggag	agcactggtt	ggettteggt	tygaacccac	gittiaacat	trantatana
21301	tttgatcctt	ttggattctc	ggatgategt	etcaaacaga	tttaccagtt	rgaacacgag
21421	ggtctcctgc	gccgcagcgc	tettgetace	aaggaccggt	gtattacgct	ggaaaaatct
21481	acccagaccg	tgcagggccc	ccgttctgcc	gcctgcggac	ttttctgctg	catgttcctt
21541	catgcctttg	tgcactggcc	tgaccgtccc	atggacggaa	accccaccat	gaaattgcta
21,501	actggagtgc	caaacaacat	gcttcattct	cctaaagtcc	agcccaccct	gtgtgacaat
21661	caaaaagcac	tctaccattt	tctcaatacc	cattcgcctt	attttcgctc	tcatcgtaca
21721	cacatcgaaa	gggccactgc	gttcgaccgt	atggatgtgc	aataatgatt	catgtaaaca
21781	acgtgttcaa	taaacagcac	tttattttt	acatgtatcg	aggctctgga	ttacttattt
21841	atttacaagt	cgaatgggtt	ctgacgagaa	tcagaatgac	ccgcaggcag	tgatacgttg
21901	cggaactgat	acttgggttg	ccacttgaat	tcgggaatca	ccaacttggg	aaccggtata
21961	tcgggcagga	tgtcactcca	cagctttctg	gtcagctgca	aagctcccag	caggtcagga
		tgaaatcaca				
		actgaaacac				
		tgcccacatc				
		tacccatggc				
		tcatcttggc				
22321	gggdcoagta	gcttgaaagc	ctactagact	ttactaccct	cootataaaa	catcccccag
22381	gacctactca	aaaactggtt	acctacacaa	ccggcatcat	tcacacagca	acaaacatca
22301	ttattaacta	tttgcaccac	acttctqccc	caggaatttt	gggtgatttt	gattcactca
22501	cogciggeda	tcaaggctcg	ttatacatta	teactageca	catccatctc	gataatctgc
22301	toottottott	tcataatatt	cagatagaa	casttgacct	tereste	atcattacaa
22201	teettetgaa	tcataatatt	gecatgeaay	taccicaget	antananah	atcattycay
22021	ccatgaggcc	acaacgcaca	geetgtaeat	teccaattat	ggrgggggar	Ctyayaaaaa
22081	gaatgtatca	ttccctgcag	aaatetteee	accategige	caguguett	greattagre
22/41	aaagttaact	ggatgcctcg	gractcerea	ttcacgtact	ggtgacagat	gegettgtat
22801	tgttcgtgct	gctcaggcat	tagtttaaaa	gaggttctaa	gcccgctatc	cageetgtae
22861	ttctccatca	gcagacacat	cacttccatg	CCTTTCTCCC	aagcagacac	caggggcaag
22921	ctaatcggat	tcttaacagt	gcaggcagca	gctcctttag	ccagagggtc	atctttggcg
		tgcttcttt				
23041			ctcttctctt	tcttcttcgc	tgtcttgact	gatgtcttgc
	cccactgcta	caagttgcgc				
23101	atggggacat	gtttggtctt	ccttggcttc	tttttcgggg	gtatcggagg	aggaggactg
23161	atggggacat tcgctccgtt	gtttggtctt ccggagacag	ccttggcttc ggaggattgt	gacgtttcgc	tcaccattac	caactgactg
23161 23221	atggggacat tcgctccgtt tcggtagaag	gtttggtctt ccggagacag aacctgaccc	ccttggcttc ggaggattgt cacacggcga	gacgtttcgc caggtgtttc	tcaccattac tcttcggggg	caactgactg cagaggtgga
23161 23221 23281	atggggacat tcgctccgtt tcggtagaag	gtttggtctt ccggagacag aacctgaccc aagggctgcg	ccttggcttc ggaggattgt cacacggcga gtccgacctg	gacgtttcgc caggtgtttc gaaggcggat	tcaccattac tcttcggggg gactggcaga	caactgactg cagaggtgga accettecg
23161 23221 23281	atggggacat tcgctccgtt tcggtagaag	gtttggtctt ccggagacag aacctgaccc	ccttggcttc ggaggattgt cacacggcga gtccgacctg	gacgtttcgc caggtgtttc gaaggcggat	tcaccattac tcttcggggg gactggcaga	caactgactg cagaggtgga accettecg

FIG. 28A-6

		aggcagagaa	acaacacaca	togaaactca	gccattgctg	tcaacatcgc
23401	gtgttctcct	atcacatctc	atactagaca	acqaqqaaaa	ggagcagagc	ttaagcattc
23461	cacgagtgcc	tcctgccacc	agetetageg	tagaagataa	ggaggtcgac	gcatctcatg
23521	caccgcccag	taaaaaagcg	acciccaccc	aggaagacat	cgaacaagaC	ccgggctatg
23581	acatgcagaa	taaaaaagcg	aaagagtetg	agccagacac	adadadag	gatgaaaact
23701	occcaaaaca	ggaacacgag gcaagcggat	aactatcacc	aagatgctgg	adatayyyac	cagaacacce
23761	actacctcat	agggcttgac	ggggaagacg	cgctccttaa	acatetagea	ayacayccac
22701	tratantraa	agggettgae ggatgeatta	ttggacagaa	ctgaagtgcc	catcagtgtc	gaagagetta
23021	~cacagecaa	ggatgcatta cgagcttaac	ctattttcac	ctcgtactcc	ccccaaacgt	cagccaaacg
23881	geegegeeta	gccaaatcct	cocttaaact	tttatccagc	ttttgctgtg	ccagaagtac
23941	gcacctgcga	tcacatcttt	tttaaaaatc	aaaaaattcc	agtctcctgc	cgcgctaatc
24001	tggctaccta	cgatgcccta	ctcaatctcc	gacctggttc	acocttacct	gatatagctt
24061	gcacccgcgc	ggttccaaag	ctcaacccgg	atctagacaa	taatgagact	cgggccgcaa
24121	ccttggaaga	ggttccaaag	accucegagg	atgaggatga	cagcatteta	gtggaattgg
24181	atgctctgca	aaagggagaa	aatggcatgg	acgageacea	cagagacaca	cactttgcat
24241	aaggcgataa	tgccagactc	gcagtactca	agegaagege	catogaccao	ttactcatta
24301	accccgctgt	caacctgccc	cctaaagtca	tgaeggeegt	tacggaccag	gagggtaaac
24421	cagtogtcag	tgatgagcag	ctaacccgat	ggctgggcac	cgacteteec	-setatatta
24/81	aagagggtcg	caagettatg	atggccgtgg	tgctggttac	cgtagaacta	gagtgtcttt
24401	aagagegeeg	taccgattca	gaaaccttgc	gcaaactcga	agagaatctg	Cactacactt
24341	ggcgccccc	ctttgtgcgg	caggcatgca	agatatctaa	cgtggaactc	accaacctgg
24601	LLayacacyy	gggtattctg	catgagaatc	gcctaggaca	aagcgtgctg	cacagcaccc
24661	Ettectacat	agcccgccgt	gattacatco	gcgattgtgt	ttatctctac	ctgtgccaca
24721	ttaaggggga	cggcatgggt	gattacaccc	aatotttaga	agaacagaac	ctgaaagagc
24781	cgtggcaaac	cggcatgggt	grarygrage	ttatataaac	aggattcaac	gagcgcaccg
24841	taaacaagct	cttacagaaa	tetettaagg	es es es es es es	cagggttact	ttocoaaacg
24901	tcgcttccga	cctacagaaa	ctcatcttcc	cagagegeee	teectette	atcctggaac
24961	gactgcctga	ctttatgagc	cagagcatgc	ttaacaattt		cctctcacct
25021	actccaatat	ctttatgage	acctgctgcg	cactgccctc	cgaccicgig	ccccccccc
25321	ctcctgggca	agattaccac	ccctatcass	tcaagttcta	tgaggaccaa	tcacagcctc
25381	. ttgccccgga	agattaccac actttcggcc	treetestes	cccadadaa	aattctggcc	caattgcaag
25/41	gaggaaaa	aggaggcaya a caagcaacag	coctaccato	: tccgctccga	, gtcgaggaac	ccggcggcgu
25861	Cocagoago	z gatgggacg-	atacaaqtco	taacaaggg	: ataagaatgo	catcatctcc attccatcat
2592	L ggtaagaag	accygcaggg	caacatatco	ttcacgcgg	gctacttgct	attccaccat ccctactat
26043	L ggggtgaac	t ttccgcgcaa	t tyttttgcat	. caccacego	acaacaacct	cccctactat ccaacagaaa taaagattac
2628	1 catcttcca	g cagagtcggg	gccaagagca	a ggaactgaa	acaaaaaac	gatetetgeg geactetega
2634	1 ttcgctcac	c agaagttgtt	: tgtatcacaa	a gagcgaagat	caacttcago	gcactctcga aggcagcgac
2640	1 adacaccas	a actetette	a acaagtacto	g cgcgctgac	. cttaaagagi	aggcagcgac a gaaattccca
2040	1 ggacgeega	t cassassagg	gggaattac	a tcatcctcg	a catgagtaaa	gaaattccca tcccaggact
2646	1 cgcgcttat	t ctacacas;	cagececaa	a toggattgg	ggcaggcgc	tcccaggact gttaatgata
2652	1 cgccttaca	- estassita	r ctcarcocc	g ggccttcta	t gatttctcga	gttaatgata c acgccccgcc
2658	1 actccaccc	g catyaatty	a stagtttt	g aacadtcad	tcttaccac	acgeceegee t ceegeteeca
2664	1 tacgcgcct	a ccgaaacca	a acacticity	g dacageous	ccaggaaag	t cccgctccca t gcaggtgcgc
2670	1 aacacctta	a tcccagaaa	tggedeged	g ccccagtge	aatgactaa	t gcaggtgcgc
2676	1 ccactgtat	t acttcctcga	a gacgcccag	g ccgaagice	a acceptant	t gcaggtgcgc a aaacgcctga
2682	1 agttagcog	g cggctccac	c ctatgtcgt	c acaggeete	y yearaarara	a aaacgcctga g cttggtctac
2688	1 tgatcagag	g ccgaggtate	c cagctcaac	g acgagtcgg	t gagetetee	g cttggtctac c cctcgtcagg
2600	1 maccamacm	g aatctttca	g attgccggc	t gcgggagat	c ttccttcac	c cctcgtcagg c gggaccgttc
2770	1 stattetas	c tttggaaag	t tegtetteg	c aaccccgct	c gggcggaat	c gggaccgttc a tctcctgggc
2/00	1 courticiga	a contttac	t coctatate	t acttcaacc	c cttctccgg	a tctcctgggc g gacggctacg
2706	_ aacctgtgg	a gyayttat	a cogaactto	g acgcgatta	g cgagtcagt	g gacggctacg c actgccgccg
2712	1 actacccgg	a cyaytttac	a ctgaactet	c teaactaca	a catctagac	c actgccgccg c ccaaggatca
2718	I attgatgto	t ggtgacgcg	y coguyotat m sactmattm	a otteateta	c ttcgaactc	c ccaaggatca
2724	1 ctttcgctg	ic tttgcccgg	y advicating	a gercareta		

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27301	ccctcaaggt	ccaacccaca	gagtgcggat	tactatcgaa	ggcaaaatac	actctcgcct
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2/361	ttccatctac	Lectedage		attacataca	accettecet	atettatata
27421	ttccatctac	tgcatttgta	accaccegg	actycacyda	agcccccgcc	attenance
27481	tactgagttt	aataaaaact	gaattaagac	teteetaegg	actgeegett	Cucaacccy
27541	gattttacaa	ccagaagaac	gaaacttttc	ctgtcgtcca	ggactctgtt	aacttcacct
27601	tteetaetea	caaactagaa	gctcaacgac	tacaccgctt	ttccagaagc	attttcccta
27661	ctaatactac	tttcaaaacc	ggaggtgagc	tccaaggtct	tcctacagaa	aacccttggg
27721	tggaagcggg	cottataata	ctaggaattc	ttacaaataa	gcttgtgatt	attetttget
2//21	acctatacac	ccctycayty	actttcctac	taatattata	gtattggttt	aaaaaataaa
2//81	acctatacac	accetycete	actecetag	esttttes.	accepted	coasttacca
27841	gcccatacta	grerrgerrg	ttttactttc	gettttggaa	ccyggccccg	ccaaccacga
27901	tccatgtcta	gacttcgacc	cagaaaactg	cacacttact	tttgcacccg	acacaageeg
27961	catctgtgga	gttcttatta	agtgcggatg	ggaatgcagg	tccgttgaaa	ttacacacaa
28021	taacaaaacc	tggaacaata	ccttatccac	cacatgggag	ccaggagttc	ccgagtggta
28081	cactgtctct	atccaagatc	ctgacggttc	catccgcatt	agtaacaaca	ctttcatttt
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20141	caaggacaac	atcataacat	tetecattee	ttattactta	tacacttacc	ttettactee
20201	tttactgtgc	attgtaatgt	coccattge	aaccactccc	atcasasacc	ccaataacaa
28261	tttactgtgc	gtatgeatae	accegation	aaccacccgc	accaaaaacg	tetetestat
28321	agaaaaaatg	ccttaacctc	tttctgttta	cagacatgge	Licitata	Lucutatat
28381	ttgtcagcat	tgtcactgcc	gctcacggac	aaacagtcgt	ctctatccct	ctaggacata
28441	attacactct	cataggaccc	ccaatcactt	cagaggtcat	ctggaccaaa	ctgggaagcg
28501	ttgattactt	tgatataatc	tgcaacaaaa	caaaaccaat	aatagtaact	tgcaacatac
28561	aaaatcttac	attoattaat	gttagcaaag	tttacagcgg	ttactattat	ggttatgaca
20501	gatacagtag	tcastataca	aattacttoo	ttcatattac	ccaqttaaaa	accacgaaaa
20021	tgccaaatat	gggaaatt	castcasta	acaattetet	agaaactttt	acatetecca
2898T	tgccaaatat	ggcaaayatt	cyattegaty	acaaccccc	agadactett	acaccccca
28/41	ccacacccga	cgaaaaaaac	accecagact	caatgattgc	aattyttyta	gcggrggcag
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28861	atcctaaaaa	acaagatctc	ctactaaggc	ttaacattta	atttctttt	atacagccat
28921	ggtttccact	accacattcc	ttatgcttac	tagtcttgca	actctgactt	ctgctcgctc
28981	acacctcact	gtaactatag	gctcaaactg	cacactaaaa	ggacctcaag	gtggtcatgt
29041	cttttggtgg	agaatatatg	acaatggatg	otttacaaaa	ccatgtgacc	aacctggtag
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29101	ctattatgga	aacggcagag	account	arattataar	attattatac	taccatctac
23191	ctattatgga	accyactata	adaytayttt	torrorat	attactycac	atagaattta
29221	cactccagca	ccccgcacaa	ctactttctc	Lageageage	gucyctaaca	atacaactic
29281	caatccaacc	tttgccgcgc	ttttaaaacg	cactgtgaat	aattctacaa	cttcacatac
29341	aacaatttcc	acttcaacaa	tcagcattat	cgctgcagtg	acaattggaa	tatctattct
29401	tgtttttacc	ataacctact	acgcctgctg	ctatagaaaa	gacaaacata	aaggtgatcc
29461	attacttaga	tttgatattt	aatttgttct	tttttttt	atttacagta	tggtgaacac
29521	caatcatggt	acctagaaat	ttcttcttca	ccatactcat	ttgtgcattt	aatgtttgcg
20581	ctactttcac	acceptance	acarcaaccc	cagactgtat	aggaggattt	acttcctata
29301	cactttttgc	ttttattaat	tagetataga	tatataacat	antetaceta	ottattaatt
29041	Cacciffication	tetegetace	cycattegey	cacgtageac	agteegeeeg	catccccaat
29701	ttttccaact	tetagaetgg	accertgige	gaattyccta	cctgcgccac	caccccgaac
29761	accgcaacca	aaatatcgcg	gcacttctta	gactcatcta	aaaccatgca	ggetataeta
29821	ccaatatttt	tgcttctatt	gcttccctac	gctgtctcaa	ccccagctgc	ctatagtact
29881	ccaccagaac	accttagaaa	atgcaaattc	caacaaccgt	ggtcatttct	tgcttgctat
29941	cgagaaaaat	cagaaattcc	cccaaattta	ataatgattg	ctggaataat	taatataatc
30001	tgttgcacca	taatttcatt	tttgatatac	cccctatttg	attttggctg	gaatgctccc
30061	aatgcacatg	atcatccaca	agacccagag	gaacacattc	ccctacaaaa	catgcaacat
30101	ccaatagcgc	tagtaratta	cussautuss	CCacaacccc	cactactece	toctattagt
20101 20101	toattagege	taacayatta	agazag cyaa	aacactcacc	acctccaatt	
POTRI	tacttcaacc	Laaccggcgg	ayacyactya	acactcact	anananatan	acatacaca
30241	tctgctcgat	atggacggcc	gcgtctcaga	acagegaett	geccaactac	geaceegeea
30301	gcagcaggaa	cgcgcggcca	aagagctcag	agatgtcatc	caaattcacc	aatgcaaaaa
30361	aggcatattc	tgtttggtaa	aacaagccaa	gatatcctac	gagatcaccg	ctactgacca
30421	tegeetetet	tacgaacttg	accccaacq	acaaaaattt	acctgcatgg	tgggaatcaa
30481	ccccatagtt	atcacccage	aaagtggaga	tactaagggt	tgcattcact	gctcctgcga.
30541	ttccatcgag	tocacctaca	ccctactass	gaccctatoc	ggcctaagag	acctoctacc
30541	aatgaattaa	aaaatratta	ataasaaata	acttacttca	aatcaccaat	aaggteteta
2000T	aaryaarida	aaaacyatta	acaaaaaacc	acctattaca	aactotoota	ttotacacca
2000T	ttgaaatttt	ccccagcag	Caccicactt		aaccctygta	gtactatact
30721	cgttcagcgg	catactttct	ccatacttta	aaggggatgt	caaattttag	GEGGEGEGE
30781	gtacccacaa	tcttcatgtc	tttcttccca	gatgaccaag	agagtccggc	tcagtgactc
30841	cttcaaccct	gtctacccct	atgaagatga	aagcacctcc	caacacccct	ttataaaccc
30901	agggtttatt	tccccaaatg	gcttcacaca	aagcccagac	ggagttctta	ctttaaaatg
30961	tttaacccca	ctaacaacca	cagggggatc	tctacaccta	aaagtgggag	ggggacttac
31001	agtggatgac	actratracta	cotteraara	aaacatacct	getacageae	Ccattactaa
31001	aaataatcac	tototacagaca	totacaaya	agatorotto	daactcaaa	acastasact
31144	aaatdatcac	theresees	-attoacty	tagasaaat	gadactcada	taaaggaaact
21,141	atgtgccaaa	LEGGGAAAEG	ygildadait	caacaacygt	yacacccyca	-aaayya tay

FIG. 28A-8

	t - t t	ttatggactg	raataaaccc	tccacctaac	tgtcaaattg '	tggaaaacac
31201	tattaacacc	gatggcaaac	ttactttact	attagtaaaa	aacggagggc	ttgttaatgg
31261	taatacaaat	gatggcaaac ctagttggtg	tatcacacac	totoaaccaa	atgttcacac	aaaagacagc
31321	ctacgtgtct	ttaagattat	otttagacac	trctggaaat	ctattaactg	atgaatcaga
31381	aaacatccaa	ccacttaaaa	atticigation	tacagegaace	agtgaaactg	tagccagcag
31501	caaagccttt	atgccaagta	ctacayetta	andtacted the contract	gatagaagtc	tatttccctt
31561	aaactacatt	catggaatat	gttactacat	gactagetae estteettee	aatottocct	atoccataca
31681	atttgaatgg	ataatgctaa	caagtgaatc	tecagaaage	taaataaaa	tttaagtgtt
31741	atcccccttt	ttcttttctt	acattacaga	agacyacaac	caccttccca	tttgacagaa
31801	tttatttaaa	atcacaaaat	tcgagtagtt	attitgeete	taccetteca	gatagacatt
31861	tacaccaatc	tctccccacg	cacagcttta	aacatttgga	at at areast a	agtgatagat
31921	gttttagatt	ccacattcca	aacagtttca	gagcgagcca	acceggggee	caastacass
31981	aaaaatccat	cgcgatagtc	ttttaaagcg	ctttcacagt	ccaactgctg	tocasasaca
32041	tccggagtct	ggatcacggt	catctggaag	aagaacgatg	ggaatcataa	ecctacatac
32101	gtatcggacg	attgtgtctc	atcaaaccca	caagcagccg	ctgtctgcgt	eteccetta
32161	aactgctgtt	tatgggatca	gggtccacag	tgtcctgaag	catgatttta	acageeeeea
22221	acatcaactt	tatgggatca	tgcgcgcagc	aacgcattct	gatttcactc	aaatctttgc
22221	acateaacco	acacattatt	acaatattgt	ttaataaacc	ataattaaaa	gcgctccagc
32201	agraggraca	acacattatt	atcocccctg	catgaccatc	ataccaaagt	ttaatataaa
32341	thantaga	ttccctcaaa	aacacactac	ccacatacat	gatctctttt	ggcatgtgca
32461	tattaacaat	cactgccaac	accoctcccc	cagccatgca	ttgaagtgaa	ccctgctgat
32521	teeggaacea	atgaagaacc	caattetete	gaccotgaat	cacttgagaa	tgaaaaatat
32581	tacaatgaca	atgaagaacc	catasatoca	tgcatcttct	cataattttt	aactcctcag
32641	ctatagtggc	catatcccag	cacaaacgea	actettacaa	aacagtaaag	ctggcagaac
32701	gatttagaaa	catateceag	gyaacaggaa	gccccagons	agtatcacaa	tctggcaaca
32761	aaggaagacc	acgaacacaa ttcagtcata	Cttacactac	tttcattttc	ctcacaacqt	ggtaactggg
32821	. gcgggtggtc	ttcagtcata	gaagereggg	atataaaaa	tocococaac	cttotcataa
32881	. ctctggtgta	agggtgatgt	ctggcgcatg	atglegageg	accordecet	ggcagaacac
32941	. tggagttgct	tcctgacatt	ctcgtatttt	gtatagtaaa	tatastaatt	caagtacage
33001	actcttcttc	gccttctatc	ctgccgctta	gegraticeg	tostassasa	tccatcgcat
33061	cacactctta	gccttctatc agttggtcaa	aagaatgctg	getteagttg	Laattaaaat	accastocas
33121	ctaattgtto	. agttggtcaa : tgaggaaatc	atccacggta	gcatatgcaa	attectaatta	ageaacgeaa
33181	ctggattgcg	tgaggaaacc tttcaagcag	gagaggagag	ggaagagacg	gaagaaccat	etetecccc
33241	attccaaacc	atctcgcagt	acttcaaatt	gtagatcgcg	cagatggcat	eretestes
22201	cactotototo	gtgaaaaagc	acagctaaat	caaaagaaat	gcgattttca	aggtgeteaa
22261	cactgogott	gtgaaaaagc caacaaagcc	tccacgcgca	catccaagaa	caaaagaata	ccaaaagaag
22421	. cggcggccc	taactcctca	atcatcatat	tacattcctg	caccattccc	agataatttt
22461	gageattect	taactcctca gccttgaatt	attcgtgtca	gttcttgtgg	taaatccaat	ccacacatta
33461	L cagettecet	ccggagggcg	ccctccacca	ccattcttaa	acacaccctc	ataatgacaa
3354.	L caaacayyco	ccggagggcg tcctgtgtca	cctgtagcga	attgagaatg	gcaacatcaa	ttgacatgcc
3360.	L aataccity	agttcttctt	taagttctag	ttgtaaaaac	: tctctcatat	tatcaccaaa
3366.	l Cttggctcte	agaagccccc	caaaaaaaa	agcagggag	gctacagtgc	agtacaagcg
3372.	L ctgcttaged	caattggctc	cadcaaaaaa	aagattggaa	taagcatatt	gggaaccgcc
3384	l agtaatatca	tttgccaaaa	. cygaaucaca	aacctctgg	atgcaaatgc	: aataggttac
3390	l aaaagaaaa	tttgccaaaa	aaacacccac	attagtctgg	aaaaataaaa	aaaaaaacaa tcacaagaca
3396	1 cgcgctgcg	tccaacatty		tocagoogs	agtetteca	tcacaagaca acaacagcac
3402	1 gcgtcatate	c atagtageet	gacyaacays	, eggacadace	tootoattaa	acaacagcac atccagacat
3408	1 agccacagg	g tetecagete	gacccccgc	. aattetteat	gaagcataca	atccagacat
3414	1 cgaaagttc	e tegeggtgae	cagcalgaa	. aactettgat	- ttaaatataa	ttatgcttaa gagaataaaa
3420	1 gttagcatc	a gttaacgaga	aaaaacagco	- abcaraget	. seegggcaca	gagaataaaa
3426	1 tcgtaagta	t agcaaagcca	cccccgcg	atacaaagta	adaggeded	gagaataaaa ctaaatacac
3432	1 aatataatt	a tttctctgct	gctgttcag	g caacgtege	- 2009	ctaaatacac cacaagctct
3438	1 atacaaagc	c tcatcagcca	tggcttacca	a gacaaagta	agogggcac	cacaagctct acgtaatggg
3444	1 aaagtcact	c tccaacctct	ccacaatat	a tatacacaa	g ccclaaact	g acgtaatggg g tcaccaggga
3450	1 agtaaagtg	t aaaaaatcc	gccaaaccc	a acacacacc	c cgaaactgc	tcaccaggga cacggtacgt
3/156	1 aaagtacag	t ttcacttcc	g caatcccaa	c aagcgtcac	r ráczerzec,	cacggtacgt tage
3420	1 cacateces	t taacttoca	cgtcatttt	c ccacggccg	c accaccca.	t ttagccgtta t tacatattgg
2460	1 200002020	c caatcaccac	acaccccac	a accettada	a tcacctcat	t tacatattgg
2171	1 caccattee	a tctataagg	t atattattg	a tgatg		
24/4	ID NO: 12	a cocacass.		-		
SEU	140. IA		_			

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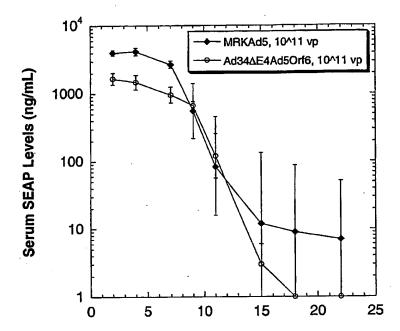


FIG. 29

			Pre Wk4 Wk8			Wk	24	Wk 28		AAK 20				
r	Vaccine	Monkey	P	re	W	(4					Mock	Gaq	Mock	Gag
-1		ID .	Mock	Gag*	Mock	Gag	Mock	Gag	Mock	Gag	MOCK	Gay	MOOK	
	Wk 0, 4, 24 MRKAd5gag, 10^11 vp MRKAd5gag, 10^11 vp MRKAd5gag, 10^11 vp Ad346E1gagAE4Ad5Ori6, 10^11 vp Ad346E1gagAE4Ad5Ori6, 10^11 vp	00C018 00C034 00C058 00D038 00D042 00D068	1 0 4 6 6	5 4 4 8 30 18	13 5 3	1025 219 1088 111 89 118	0 5 0	824 404 440 301 264 816	8 3 4 0 1	756 445 1439 224 73 429	0 3 0	474 339 2338 536 181 439	0 0 0	383 216 940 233 69 273
١	Ad34AE1gagAE4Ad5Orl6, 10411 vp	ODDOG	ľ						<u> </u>		<u> </u>			

Vaccine	Monk ID	•	D4 ⁺ CD3 ⁺ mphocytes	IFN-γ ⁺ CD8 ⁺ CD3 ⁺ per 10 ⁶ Lymphocytes		
	Ī	Mock	Gag ^a	Mock	Gag ^a	
Ad34ΔE1gagΔE4Ad5Orf6	00D038	22	154	130	450	
	00D042	32	118	96	171	
	00D066	12	238	150	442	

		Monkey		70	Test	wks	T≕e	wke	T=24	wka_	T=20	WKB	1=32	WKS
Vaccine	Vaccine			Goo	Mock	Ged	Mock	Gag	Mock	Gas	Mock	Gag	Mock	Gag
T=0, 4 wke	T=24 w/ks	ID_	Mock	Cay	MOCK	_	-	334	-	99	0	305	3	244
Ad34AE1gagAE4Ad5Od6, 10^11 vp	Ad35AE1gagAE4Ad5OrlB, 10^10 vp	00D016	4	8	1	84				136		493	lı	259
Ad34AE1gagAE4Ad5Od8, 10*11 vp	Ad35AE1gagAE4Ad5Orf6, 10^10 vp	000044	1 1	١,	a a	79		374	l °	145	,	351	1 1	235
Ad34AE1gagAE4Ad5Od6, 10^11 vp	Ad35AE1gagAE4Ad5OdB, 10^10 vp	00D064	1 4	6	1	125	B	655	l °	140	١ ١	١		1 [
Addate I gargate Add Cities, IV-11 VP			<u> </u>	ļ	<u> </u>		 _	54	١.	 	5	5	3	0
Naive		000087	1_1	11	3	<u> 3</u>	L	_ 04	1			<u> </u>		

Vaccine (T≔0, 4 Wks)	Vaccine (T≃24 Wk)	Monkey		D4*CD3* mphocytes	IFN-γ*CD8*CD3* per 10 ⁸ Lymphocytes		
		ID	Mock	Gag	Mock	Gag	
Ad34AE1gagAE4Ad5Orf6, 10^11 vp	Ad35ΔE1gagΔE4Ad5Orf6, 10^10 vp	00D016	62	433	176	1288	
Ad34∆E1gag∆E4Ad5Orf6, 10^11 vp	Ad35∆E1gag∆E4Ad5Orf6, 10^10 vp	00D044	136	593	323	1871	
Ad34ΔE1gagΔE4Ad5Orf6, 10^11 vp	Ad35ΔE1gagΔE4Ad5Orf6, 10^10 vp	00D064	188	785	292	892	

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